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992 amino acids
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                         MOLECULE TYPE: Protein
                                             COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                             amino acid
                                                                                                                                                                                                                                                                                                                                                                            linear
   New York
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CNGFROMN----LKDTKTLRVAKAIYKRYI-ENNSVVSKQLKPATKTYIRDGIKKQQIGSV 169
                                                             MFDQAQTEIQAVMEENAYQVFLTSDIYLEYVRSGGENTAYMS--NGGLGSLKVLCGYLPT 227
                                                                                                                          LNEEEEWTC-ADLKCKLS----PTVVGLSSKTLRATASVRSTETAE----NGFRSFKRSD 278
                                                                                                                                                                                                                                                                                                                                                            ---SYEE 433
                                                                                                                                                                                                                                                                                                                                                                            434 DPQTILDDHLSRVLKTPGCQSPGVGRYSPRSRSPDHHHQHHHHQQCHTLLSTGGKLPPVA 493
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    494 ACPLLGGKSFLTKQTT-----KHVHHHYIHHAVPKTKEEIEAEATQRVRCLCPG 543
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             EMPGSP----EDVERNOKILOWIIEGEKEISRHKKTNHGSSGVKKOLSHDMVRPLSIERP 697
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      VAVHPWV--SAQLRNVVQPSHPFIQDPTMPPNPAPNPLTQLEEARRRLEEEEKRAGKLPL 755
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STREET: 1185 Avenue of the Americas
CITY: New York
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APPLICANT: Conetentini, Franklin
APPLICANT: Cang, Li
TITLE OF INVENTION: AXIN GENE AN
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Patent No. 6307019
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CORRESPONDENCE ADDRESS:
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US-08-890-865A-1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  286 GSGYVFAPATSANDSE---LSSDALTDDSMSMTDSSVDGVPPYRMGSKKQLQREMHRSVK 342
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STWEENTYPSFLKSDIYLEYTRTGSESPKVCSDQSGGGGGGGGGKGMSGYLPTLNEDEEWKCD
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ----ADLKCKLSPTVVGLSSKTL-----RATASVRSTETAENGFRSFKRSDPVNPYHV
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380 QDADEDDGRDPLPPS--RLTQKLLLETAAPRAPSSRRYNEGRELRYGSWR--EPVNPYYV
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             SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/890,865A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
37.2%; Score 1001, 2007, 2007
Best Local Similarity 42.7%; Pred. No. 1e-139;
Matches 390; Conservative 133; Mismatches 246;
                                                                                                                                                                                                                                        0575/54249
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                              NAME: White, John P
REGISTRATION NUMBER: 28,678
REFERENCE/DOCKET NUMBER: 05.7
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212)278-0400
                                                                                                  FILING DATE: 10-JUL-1997
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
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Sequence 1, Appli
                                                                                          April 20, 2006, 15:30:33 ; Search time 70.1847 Seconds (without alignments) 989.497 Million cell updates/sec
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            GenCore version 5.1.7
Copyright (c) 1993 - 2006 Biocceleration Ltd.
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1. / GgTZ 6/ptodate/1/iaa/5_COMB.pep:*

2. / GgTZ 6/ptodate/1/iaa/6_COMB.pep:*

3. / GgTZ 6/ptodate/1/iaa/H_COMB.pep:*

4. / GgTZ 6/ptodate/1/iaa/H_COMB.pep:*

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Maximum Match 100%
Listing first 45 summaries
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	166 3.7 196 2 US-09-702-705-339 166 3.7 196 2 US-09-736-457-339 166 3.7 196 2 US-09-614-124B-339 166 3.7 196 2 US-09-589-184-339 166 3.7 196 2 US-09-589-184-339 166 3.7 196 2 US-09-658-824-339 166 3.7 196 2 US-09-658-824-339 166 3.7 196 2 US-09-651-563-339 166 3.7 196 2 US-09-61-754-339 167 3.7 196 2 US-09-61-754-339 167 3.7 196 2 US-09-61-642-339 167 3.7 196 2 US-09-941-886-21 167 3 3.7 182 2 US-09-941-886-21 160 5 3.6 119 2 US-09-865A-18 159 3.6 119 1 US-08-890-865A-18 159 3.6 181 2 US-09-103-25 159 3.6 181 2 US-09-103-25 159 3.6 181 2 US-09-103-25 159 3.6 181 2 US-09-103-25	ALIGNMENTS	Sequence 10, Application US/08890865A Sequence 10, Application US/08890865A Patent No. 6307019 GENERAL INPORMATION: APPLICANT: Constantini, Franklin APPLICANT: Sequences: 23 CORRESPONDENCES: 23 CORRESPEE: 1105 Avenue of the Americas CITY: New York CONTRY: New York COUNTRY: US ZIP: 10036 COMPUTER: New York COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: Ploppy disk COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: 10-JUL-1997 CLASSIFICATION: UNDER: 28,678 FILING DATE: 10-JUL-1997 CLASSIFICATION NUMBER: 28,678 REFERENCE/DOCKET NUMBER: 28,678 REFERENCE/DOCKET NUMBER: 28,678 REFERENCE/OCKET NUMBER: 28,678 REFERENCE/OCKET NUMBER: 28,678 REFERENCE/OCKET NUMBER: 28,678 TELECOMMUNICATION INFORMATION: TELEPHONE: (212)394-0526 INFORMATION FOR SEQ ID NO: 10: SEQUENCE CHARACTERISTICS: LENGTH: BSS amino acide TYPE: amino acid STRANDEDNESS: single TOOLOGY: linear MOLECULE TYPE: Protein	tch 39.1%; Score 1736; DB 2; al Similarity 44.5%; Pred. No. 4e-147; 406; Conservative 128; Mismatches 214	12 DPSSSPREDAPRPPVPGEEGETPPCOPSVG	62LG-EPEGRASPDSPLTRWTKSLH

Sequence Sequence Sequence Sequence Sequence

-08-890-865A-11 -08-588-258B-31

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CNGFRQMNLKDT----KTLRVAKAIYKRYI-ENNSVVSKQLKPATKTYIRDGIKKQQIGSV 169
                                                                                     -----ADLKCKLSPTVVGLSSKTLRATASVRSTETAENGFRSFKR 276
                                                                                                                                                                                                                                                             SDPVNPYHVGSGYVFAPATSANDSE---LSSDALTDDSMSMTDSSVDGVPPYRMGSKKQL 333
                                                                                                                                                                                                                                                                                  OREMHRSVKANGOVSLPHFPRTHRLPKEMTPVEPAAFAAELISRLEKLKLELESRHSLEE 393
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                                                                                                                                                  218 MFDQAQTEIQATWEENTYPSFLKSDIYLEYTRTGSESPKVCSDQSGGGTGKGISGYLPT
                                                                                                                                                                                                                 394 RLQQIREDEEKEGSEQALSSRDGAPVQ-----HPLALLPS------G
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VFEEVREDEAVLPVFEEKIIGKVEKVD 900
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STREET: 1185 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              patent No. 6307019

patent No. 6307019

GENERAL INFORMATION:

APPLICANT: Constantini, Franklin

APPLICANT: Zeng, Li

TITLE OF INVENTION: AXIN GENE AND USI

NUMBER OF SEQUENCES: 23

CORRESPONDENCE ADDRESS:

ADDRESSEE: Cooper & Dunham LLP
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                                                                --EEGDRSQDVWQWMLESERQ---SKSKPHSAQSIRKSYPLESARAAPGERVSRHHLLCA
                                                                                                              SGHSRSVARAHPFTQDPAMPPLTPPNTLAQLEEACRRLAEVSK-----PQKQRCCVASQQ
                                                                                                                                           807 AQLRNSVQPSHLFIQDPTMPPNPAPNPLTQLEBARRLEBEBEKRANKLPSKQRYVQAVMQ
                                                                                                                                                                             RDRNHSAAGOAGASPFANP-----SLAPEDHKEPKKLASVHALQASELVVTYFF
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DPSSSFREDAPRPPVPGEEGETPPCQPSVGKVQSTKPMPVS--
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CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: White, John P
RECISTRATION NUMBER: 28,678
REFERENCE/DOCKET NUMBER: 0575/54249
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 4, Application US/08890865A; Patent No. 6307019; GENERAL INFORMATION:
APPLICANT: Constantini, Franklin; TITLE OF INVENTION: AXIN GENE AND USF; NUMBER OF SEQUENCES: 23
CORRESPONDENCE ADDRESS:
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/890,865A
FILING DATE: 10-JUL-1997
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980 FEEKIIGKVEKVD 992
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INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
LENGTH: 900 amino acids
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
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STRANDEDNESS: si
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4.7%; Score 208; DB 2;
Best Local Similarity 30.8%; Pred. No. 3.1e-10;
Matches 56; Conservative 34; Mismatches 58
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Patent No. 6830914
GENERAL INFORMATION:
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US-09-498-959-2
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LENGTH: 235
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30.8%; Pred. No. 3.1e-10;
.ive 34; Mismatches 58; Indels 34; Gaps
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Patent No. 6274362

GRNREAL INFORMATION:

APPLICANT: Hodge, Martin R.

APPLICANT: Yowe, David

TITLE OF INFORTION: RGS-Containing Molecules and Uses Thereof
FILE REPERENCE: S800-19, 035800/174680

CURRENT FILING DATE: 1999-02-04

NUMBER OF SEQ ID NOS: 4

SOFTWARE: PatentIn Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 127;
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                                                                                                                       OPERATING SYSTEM: PC_DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                       0575/54249
                                                                                                                                                                         APPLICATION NUMBER: US/08/890,865A FILING DATE: 10-UUL-1997 CLASSIFICATION: 435
                                                                                                                                                                                                                        CLASSIFICATION: 4.3.3
ATTORNEY/AGENT INFORMATION:
NAME: White, John P
REGISTRATION NUMBER: 28,678
REFERENCE/DOCKET NUMBER: 0575,
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212)278-0400
                                                                                       E: Floppy disk
IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                         INFORMATION FOR SEQ ID NO: 19: SEQUENCE CHARACTERISTICS: LENGTH: 127 amino acids TYPE: amino acid STRANDEDNESS: single
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Best Local Similarity 30.8%
Matches 56; Conservative
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                                                      ZIP: 10036
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
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     New York
New York
Y: US
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   CITY: Ne.
STATE: N.
COUNTRY:
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US-09-244-314-2
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APPLICANT: Hodge David
TITLE OF INVENTION: RGS-Containing Molecules and Uses Thereof
FILE REFERENCE: 5800-19, 035800/174680
CURRENT APPLICATION NUMBER: US/09/894,749
CURRENT FILING DATE: 2001-06-27
PRIOR APPLICATION NUMBER: 09/244,314
PRIOR APPLICATION NUMBER: 09/244,314
PRIOR PILING DATE: 1999-02-04
NUMBER OF SEQ ID NOS: 4
SOFTWARE: Patentin Ver. 2.0
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Patent No. 6410240
| Patent No. 6410240
| GRUERAL INFORMATION:
| APPLICANT: Howe, David
| APPLICANT: Yowe, David
| TITLE OF INVENTION: RGS-Containing Molecules and Uses:
| TITLE OF INVENTION: RGS-Containing Molecules and Uses:
| TITLE OF INVENTION: Thereof:
| FILE REFERENCE: 5800-19A
| CURRENT FILION DAFE: 2000-02-04
| EARLIER APPLICATION NUMBER: 09/244,314
| RARLIER PILING DATE: 1999-02-04
| NUMBER OF SEQ. ID NOS: 12
| SOPTWARE: FastSEQ for Windows Version 3.0
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                                          CGEEI PYRRMLKAQSLTLGHFKEQLSKKGNYRYYFKKASDEFACGAVFEEIWDDETVLPM 827
                                                                  83 HSLLGDQDGAYLFRTFLEREKCVDTLDFWFACNGFRQMNLKDTKTLRVAKAIYKRYIENN 142
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               88 DKLLSHRDGVDAFTRFLKTEFSEENIEFWVACEDFKKCK-EPQQIILKAKAIYEKFIQND 146
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  143 SVVSKQLKPATKTYIRDGIKKQQIGSVMFDQAQTEIQAVMEENAYQVFLTSDIYLEYV 200
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APPLICANT: Wowe, David
TITLE OF INVENTION: RGS-Containing Molecules and Uses Thereof
FILE REFERENCE: 5800-19, 035800/174680
CURRENT APPLICATION NUMBER: US/09/244,314
CURRENT FILING DATE: 1999-02-04
NUMBER OF SEQ ID NOS: 4
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 4
LENGTH: 235
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APPLICANT: Fowe, David
TITLE OF INVENTION: RGS-Containing Molecules and Uses
TITLE OF INVENTION: Thereof
FILE REFERENCE: 5800-19A
CURRENT APPLICATION NUMBER: US/09/498,959
CURRENT FILING DATE: 2000-02-04
EARLIER APPLICATION NUMBER: 09/244,314
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; Pred. No. 6.9e-09;
37; .Mismatches 69
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ilarity 28.1%; Pred. No. 6.9e-09;
Conservative 37; Mismatches 69
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    194 AQPPMQHLSQQPLASFSSSSSGGSISL--
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Patent No. 6274362
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Patent No. 6410240
GENERAL INFORMATION:
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                                                                                                                       828 YEGRILGKVERID 840
                                                                                                                                                   301 FGDKAMGLVKPSD 313
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SOFTWARE: FastSEQ for Wi
SEQ ID NO 4
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Best Local Similarity
Then 50; Conserve
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TYPE: PRT
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                                                                                                                                                                                                                                                                                                                                                      361 EMTPVEPAAFAAELISRLEKLKLELESRHSLEERLQQ-----IREDEEKEGSEQALSSR 414
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         415 DGAPVQHPLALLPSGSYEEDPQTILDDHLSRVLKTPGCQSPGVGRYSPRSRSPDHHHQHH 474
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                                                                                                                                                              41 GKVQSTKPMPVSSNARRN-----ED-----GLGEPEGRASPDSPLTRWTKSL 82
                                                                                                                                                                                      29 GKEETSKEAKIRAKERNRLSLLVQKPEFHEDTRSSRSGHLAKETRVSPEEAV-KWGESF 87
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Homburger et al.
TITLE OF INVENTION: Nucleic acids and proteins of Drosophila melanogaster
TITLE OF INVENTION: Nucleic acids and proteins of Drosophila melanogaster
CURRENT FILE REFERENCE: File Reference: 7326-094
CURRENT FILING DATE: 1999-03-17
NUMBER OF SEQ ID NOS: 62517
SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     595 GGMSSAAGGPQLPGEEGDRSQDVWQWMLESERQSKSKP-HSAQSIRKSYPLESARAAPGE
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20.9%; Pred. No. 5.9e-09;
tive 64; Mismatches 130; Indels 196;
                                                                               Length 235
                                                                                                                       58; Indels
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                                                                                                                     34; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; Sequence 43189, Application US/09270767; Patent No. 6703491; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; ORGANISM: Drosophila melanogaster US-09-270-767-43189
                                                                        Query Match
Best Local Similarity 30.8°
Matches 56; Conservative
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Best Local Similarity 20.99
Matches 103; Conservative
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TYPE: PRT
ORGANISM: Homo sapiens
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LD 200
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US-09-270-767-43189
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                                      US-09-894-749-2
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                                                                                                88 DKLLSHRDGVDAFTRFLKTEFSEENIEFWVACEDFKKCK-EPQQIILKAKAIYEKFIQND 146
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-PEGRASPDSPLTRWTKSL 82
                    29 GKEETSIEAKIRAKEKRNRLSLLLQRPDFHGETQASRSALLAKETRVSPEEAV-KWAESF 87
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                                                                                                                                                  143 SVVSKQLKPATKTYIKDGIKKQOIGSVMFDQAQTBIQAVMEBNAYQVFLTSDIYLEYV 200
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                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: RGS-Containing Molecules and Uses Thereof FILE REFRENCE: 5800-19, 035800/174680
CURENT APPLICATION UNBER: 12/09/894,749
CURRENT FILING DATE: 2001-06-27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 235;
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Mismatches 69;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    4.3%; Score 193;
28.1%; Pred. No. 6
:ive 37; Mismatch
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1185 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-08-890-865A-23; Sequence 23, Application US/08890865A; Sequence 23, Application US/08890865A; Patent No. 6307019; GENERAL INFORMATION: APPLICANT: Constantini, Franklin APPLICANT: Zeng, Li TITLE OF INVENTION: AXIN GENE AND US; CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT FILING DATE: 2001-06-27
PRIOR APPLICATION NUMBER: 09/244,314
PRIOR FILING DATE: 1999-02-04
SUPMER OF SEQ ID NOS: 4
SOFTWARE: PatentIn Ver. 2.0
                                                                                                                                                                                                                                                                              Sequence 4, Application US/09894749
Patent No. 6830914
GENERAL INFORMATION:
APPLICANT: Hodge, Martin R.
APPLICANT: Yowe, David
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
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Matches 50; Conservative
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SOFTWARE: Patenti
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STATE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               361 RRNESPGAPPAGKADKMMKSFKPTSEEALKWGESLEKLLVHKYGLAVFOAFLRTEFSEEN
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TITLE OF INVENTION: BEPHRIN REGULATION OF G-PROTEIN COUPLED
TITLE OF INVENTION: CHEMOATTRACTION
FILE REFERENCE: 2535/106
CURRENT APPLICATION NUMBER: US/10/113,794A
CURRENT FILING DATE: 2002-04-01
NUMBER OF SEQ ID NOS: 6
SOFTWARE: FASTSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                      Length 51;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 189.5; DB 2;
Pred. No. 5.3e-08;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              168 SVMFDQAQTEIQAVMBENAYQVFLTSDIYLEYV 200
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                                                                                                                                                                                                                                                                                                                                                                        9; Mismatches
                                                                                                                                                                                                                                                                                                                                         Score 190;
Pred. No. 1
CLASSIFICATION: 435
TORNEY/AGENT TWO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 2, Application US/10113794A Patent No. 6919313
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              26;
                                                                    ATTORNEY/AGENT INFORMATION:
NAME: White, John P
REGISTRATION NUMBER: 28,678
                                                                                                                   REFERENCE/DOCKET NUMBER: 05
TELECOMMUNICATION INFORMATION: (12.2)278-0400
TELEPHONE: (212)391-0526
                                                                                                                                                                                       INFORMATION FOR SEQ ID NO: 23: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             4.3%;
                                                                                                                                                                                                                                                                                                                                         4.3%;
                                                                                                                                                                                                                        LENGTH: 51 amino acids
                                                                                                                                                                                                                                                                                                                                                       Best Local Similarity 66.7
Matches 34; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                           single
                                                                                                                                                                                                                                                                         TOPOLOGY: linear MOLECULE TYPE: Protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: Homo sapiens
                                                                                                                                                                                                                                         TYPE: amino acid
STRANDEDNESS: si
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Best Local Similarity
Matches 49; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                        US-08-890-865A-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-10-113-794A-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-10-113-794A-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQ ID NO 2
LENGTH: 519
                                                                                                                                                                                                                                                                                                                                         Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: PRT
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56 RRNEDGLGEPEGRA----SPDS-PLTRWTKSLHSLLGDQDGAYLFRTFLEREKCVDT 107
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               108 LDFWFACNGFROMNLKDTKTLRVAKAIYKRYIENNSVVSKOLKPATKTYIRDGIKKOOIG 167
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 91 GAYLFRTFLEREKCVDTLDFWFACNGFR@MNLKDTKTLRVAKAIYKRYIENNSVVSKQLK 150
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                815 GLEVFQAFLRTEFSEENLEFWLACEDFKKVK-SQSKMAAKAKKIFABFIAIQACKEVNLD 873
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       32 ETPPCOPSVGKVOSTKPM-PVSSNARRNEDGLGEPEGRASPDSPLTRWTKSLHSLLGDQD 90
                                                                                                                                                                                                                                                                                                                               4.3%; Score 189.5; DB 2; Length 520;
32.0%; Pred. No. 5.3e-08;
iive 26; Mismatches 67; Indels 11; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  24; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       151 PATKTYIRDGIKKQQIGSVMFDQAQTEIQAVMEENAYQVFLTSDIYLEYV 200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 4.0%; Score 180; DB 2; Length 930; Best Local Similarity 28.2%; Pred. No. 1e-06; Matches 48; Conservative 31; Mismatches 67; Indels 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 1, Application US/10113794A
Patent No. 6919313
GENERAL INFORMATION:
APPLICANT: FLANGAN et al.
TITLE OF INVENTION: B EPHRIN REGULATION OF G-PROTEIN COUPLED
TITLE REPERENCE: 25354106
CURRENT FILING DATE: 2002-04-01
NUMBER OF SEQ ID NOS: 6
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   168 SVMFDQAQTEIQAVMEENAYQVFLTSDIYLEYV 200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           479 RGCFDLAQKRIFGLMEKDSYPRFLRSDLYLDLI 511
PRIOR APPLICATION NUMBER: 08/09/949,016

PRIOR APPLICATION NUMBER: 60/241,755

PRIOR FILING DATE: 2000-10-20

PRIOR FILING DATE: 2000-10-3

PRIOR FILING DATE: 2000-10-03

PRIOR FILING DATE: 2000-09-08

NUMBER OF SEQ ID NOS: 207012

SOUTWARE: PastSEQ for Windows Version 4.0

SEQ ID NO 9918

LENGTH: 520
                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity 32.0
Matches 49; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; SEQ ID NO 1

: LENGTH: 930

: TYPE: PRT

; ORGANISM: Wus musculus

US-10-113-794A-1
                                                                                                                                                                                                                                             ; TYPE: PRT
; ORGANISM: Human
US-09-949-016-9918
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 15
US-10-113-794A-1
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Search completed: April 20, 2006, 15:32:28 Job time : 71.1847 secs

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GenCore version 5.1.7
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1 MSSAVLVTLLPDPSSSFRED......BETVLPMYEGRILGKVERID 840 Gapop 10.0 , Gapext 0.5 **BLOSUM62** Scoring table: Sequence:

Total number of hits satisfying chosen parameters:

1867569 segs, 417829326 residues

Searched:

length: 0 length: 2000000000 Minimum DB seq Maximum DB seq

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Published Applications AA\_Main:\*

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2. /cgn2 6/ptodata/l/pubpaA/US08

3. /cgn2 6/ptodata/l/pubpaA/US10A

4. /cgn2 6/ptodata/l/pubpaA/US10A

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5. /cgn2 6/ptodata/l/pubpaA/US10B

7. /cgn2 6/ptodata/l/pubpaA/US10B Database :

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

Result Query  No. Score Match Length DB ID  1 4027.5 90.6 843 5 US-10-723-860-1797 Sequence 1797, 2 4027.5 90.6 843 5 US-10-723-860-1797 Sequence 1797, 2 4027.5 90.6 843 5 US-10-723-860-1797 Sequence 1797, 2 4027.5 90.6 843 5 US-10-736-116 Sequence 1797, 2 4027.5 90.6 843 5 US-10-736-116 Sequence 270, 3 1654.5 35.3 90.6 4 US-10-786-720-35 Sequence 270, 3 1612.5 36.3 90.6 4 US-10-640-92846 Sequence 270, 3 1605 36.1 900 4 US-10-640-92846 Sequence 270, 3 1605 36.1 900 4 US-10-24-049-2846 Sequence 91, Ag 11 1605 36.1 900 5 US-10-477-2384-670 Sequence 670, 3 16 16 16 16 16 16 16 16 16 16 16 16 16			de	•		SUMMARIES	
1 4027.5 90.6 843 5 US-10-723-860-1797 Sequence 175 4027.5 90.6 843 5 US-10-751-756-116 Sequence 175 165.1 5 37.2 842 3 US-10-751-756-116 Sequence 175 1612.5 36.3 91.2 4 US-10-786-720-35 Sequence 276 1612.5 36.3 92.4 4 US-10-786-728.5 Sequence 276 1612.5 36.3 92.4 4 US-10-786-728.5 Sequence 276 1612.5 36.3 92.4 4 US-10-786-728.5 Sequence 276 1605 36.1 900 4 US-10-182-936.9 Sequence 91, 1605 36.1 900 4 US-10-182-936.9 Sequence 91, 1605 36.1 900 5 US-10-182-936.9 Sequence 676 1605 36.1 900 5 US-10-80-80.8 Sequence 676 1605 36.1 900 5 US-10-80-80.8 Sequence 676 1605 36.1 900 5 US-10-97-43-30.9 Sequence 676 1605 36.1 900 5 US-10-97-43-30.9 Sequence 20, 208 4.7 235 4 US-10-99-422-13 Sequence 17, 208 4.7 235 4 US-10-99-422-13 Sequence 17, 208 4.7 235 5 US-10-99-422-13 Sequence 17, 209 84-749-149-4 Sequence 17, 209 84-749-149-4 Sequence 17, 209 84-749-149-4 Sequence 17, 209 84-749-1650 Sequence 17, 209 84-749-1650 Sequence 17, 209 84-73 3 US-09-89-422-13 Sequence 17, 209 84-749-1650 Sequence 16, 209 84-7	Result No.	Score	Query	Length	DB	ID	Description
2 4027.5 90.6 843 5 US-10-751-756-116 Sequence 118 1654.5 37.2 842 3 US-09-781-781-8 Sequence 118 1654.5 37.2 842 3 US-09-788-718-8 Sequence 118 1619.5 36.4 862 4 US-10-786-720-35 Sequence 276 1615.5 36.3 347 4 US-10-052-900A-270 Sequence 276 1615.5 36.3 847 4 US-10-264-049-2846 Sequence 276 1605 36.1 900 4 US-10-786-720-36 Sequence 276 1605 36.1 900 4 US-10-182-936A-91 Sequence 91, 1605 36.1 900 5 US-10-182-936A-91 Sequence 91, 1605 36.1 900 5 US-10-182-936A-91 Sequence 676 1605 36.1 900 5 US-10-182-936A-91 Sequence 676 1605 36.1 900 5 US-10-80-287A-670 Sequence 676 1605 36.1 900 5 US-10-989-422-3 Sequence 50, 1605 36.1 900 5 US-10-989-422-13 Sequence 20, 198.5 4.7 235 4 US-10-989-422-13 Sequence 11, 198.5 4.5 1059 5 US-10-989-422-13 Sequence 11, 198.5 4.3 235 3 US-09-894-422-13 Sequence 14, 193 4.3 235 5 US-10-989-054-4 Sequence 4, 193 5 19 4 US-10-989-054-4 Sequence 676 189.5 4.3 519 4 US-10-194-149-1 Sequence 165 189.5 4.3 519 4 US-10-989-054-4 Sequence 165 189.5 4.3 519 4 US-10-989-054-4 Sequence 165 189.5 4.3 519 4 US-10-194-149-1 Sequence 165 189.5 4.3 519 4 US-10-194-1		027.	90.6	843	ß	-10-723-860-17	; –
3 1654.5         37.2         842         3 US-09-798-811-8         Sequence 35, 1619.5           4 1619.5         36.4         982         4 US-10-092-9846         Sequence 36, 3 Sequence 31, 3 Sequence 36, 3 Sequence 31, 3 Seque	7		90.6	843	2	7	_
4 1619.5 36.4 962 4 US-10-786-720-35 Sequence 35, 1617. 36.4 912. 4 US-10-796-720-35 Sequence 277 1612.5 36.3 347 4 US-10-264-049-2846 Sequence 276 21612.5 36.3 347 4 US-10-264-049-2846 Sequence 276 21605 36.1 900 4 US-10-374-979-91 Sequence 36, 21605 36.1 900 4 US-10-374-979-91 Sequence 677 1605 36.1 900 5 US-10-477-238A-670 Sequence 677 21605 36.1 900 5 US-10-477-173-670 Sequence 677 21605 36.1 900 5 US-10-477-173-670 Sequence 677 21605 36.1 900 5 US-10-98-5828 Sequence 677 21605 36.1 900 5 US-10-98-5828 Sequence 677 218 47 225 4 US-10-09-144-3015 Sequence 27 208 4.7 235 3 US-09-894-749-2 Sequence 27 208 4.7 235 3 US-09-894-749-2 Sequence 27 218 218 218 218 219 219 219 219 219 219 219 219 219 219	m	1654.5	37.2	842	ო	8	
5         1617         36.4         912         4 US-10-092-900A-270         Sequence 284           6         1615.5         36.3         347         4 US-10-264-049-2846         Sequence 284           1         1605         36.1         900         4 US-10-786-720-36         Sequence 91,           9         1605         36.1         900         4 US-10-786-720-36         Sequence 91,           1         1605         36.1         900         4 US-10-374-979-91         Sequence 91,           1         1605         36.1         900         5 US-10-477-284-670         Sequence 91,           2         1605         36.1         900         5 US-10-477-178-670         Sequence 670           3         461         4 US-10-477-178-670         Sequence 670         Sequence 670           4         173         16.5         15.2         4 US-10-106-690-584         Sequence 171           4         7.3         16.5         15.2         4 US-10-106-691-584         Sequence 171           5         16.5         15.2         4 US-10-106-691-584         Sequence 20,           5         16.5         15.2         4 US-10-344-2         Sequence 20,           5         4.7         2.3<	4	1619.5	36.4	862	4	-786-720-3	
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1612.5   36.3   826   4 US-10-786-720-36   Sequence 36, 1605   36.1   900   4 US-10-134-991   Sequence 91, 1605   36.1   900   4 US-10-182-936A-91   Sequence 91, 1605   36.1   900   5 US-10-477-238A-670   Sequence 670   1605   36.1   900   5 US-10-477-238A-670   Sequence 670   3 0.1   900   5 US-10-477-238A-670   Sequence 670   3 0.1   900   5 US-10-477-173-670   Sequence 670   3 0.1   900   5 US-10-477-173-670   Sequence 670   3 0.1   900   5 US-10-470-173-670   Sequence 670   3 0.1   900   5 US-10-97-143-3015   Sequence 670   9 0.1	v	1615.5	36.3	347	4	US-10-264-049-2846	
1605   36.1   900   4 US-10-374-979-91   Sequence 91, 91605   36.1   900   4 US-10-182-9316A-91   Sequence 91, 91605   36.1   900   5 US-10-182-9316A-91   Sequence 670   1605   36.1   900   5 US-10-477-238A-670   Sequence 670   20 US-10-477-17-670   Sequence 670   20 US-10-477-17-670   Sequence 670   20 US-10-284-570   Sequence 670   20 US-10-284-5828   Sequence 670   20 US-10-284-2828   Sequence 70 US-10-284-3015   Sequence 70 US-10-384-3015   Sequence 70 US-10-384-3015   Sequence 70 US-10-384-3015   Sequence 11 US-10-384-3015   Sequence 12 US-10-384-3015	7 .	1612.5	36.3	826	4	US-10-786-720-36	
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1605 36.1 900 5 US-10-477-218A-670 Sequence 670 Sequence	0	1605	36.1	900	4	10-182-936A-	
1 1605 36.1 900 5 US-10-680-287A-670 Sequence 670 2	10	1605	36.1	900	Ŋ	-10-477-238A	
2 1605 36.1 900 5 US-10-477-173-670 Sequence 670 8 670 19:3 461 4 US-10-186-720-34 Sequence 670 19:3 461 4 US-10-186-720-34 Sequence 134, 131 16.5 15.5 15.4 US-10-106-699-5829 Sequence 134, 131 16.5 15.5 10.3 745 6 US-11-097-143-3015 Sequence 301 10.0 10.0 10.0 10.0 10.0 10.0 10.0	11	1605	36.1	900	Ŋ	10-680-287A	
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4 733 16.5 155 4 US-10-6699-5828 Sequence 582 2695.5 10.3 745 6 US-11-097-143-3015 Sequence 582 2692.6 4.7 2.7 2.8 4.7 2.9 84.7 2.5 3 US-09-867-550-848 Sequence 2, 208 4.7 2.3 5 3 US-09-894-749-2 Sequence 2, 208 4.7 2.3 5 4 US-10-258-311B-20 Sequence 2, 208 4.7 2.3 5 4 US-10-258-311B-20 Sequence 2, 208 4.7 2.3 5 4 US-10-989-422-13 Sequence 1, 2198.5 4.5 1059 5 US-10-899-422-13 Sequence 1, 2198.5 4.5 1059 5 US-10-899-422-13 Sequence 1, 2193 4.3 2.3 5 US-10-989-054-4 Sequence 4, 2190.5 4.3 2.3 5 US-10-989-054-4 Sequence 4, 2190.5 4.3 2.3 5 US-10-98-054-4 Sequence 4, 2190.5 4.3 2.3 4 US-10-194-749-1 Sequence 2, 2190.5 4.3 2.9 4 US-10-194-749-1 Sequence 2, 2190.5 4.3 2.9 4 US-10-194-749-1 Sequence 2, 2190.5 4.3 2.9 4 US-10-13-794A-2 Sequence 2, 2190.5 4.3 2.9 4 US-10-1428-487-14 Sequence 2, 2190.5 4.3 2.9 4 US-10-428-487-14 Sequence 2, 2190.5 4.3 2.9 4 US-10-428-487-14	13	860	19.3	461	4		
5         457.5         10.3         745         6         US-11-097-143-3015         Sequence 301           7         208         4.7         227         3         US-09-867-550-648         Sequence 84           8         208         4.7         235         3         US-09-894-22         Sequence 20,           9         208         4.7         235         4         US-10-258-311B-20         Sequence 20,           1         208         4.7         235         4         US-10-899-022         Sequence 20,           1         198.5         4.5         105         5         US-10-899-422-13         Sequence 11,           2         4.5         105         5         US-10-899-422-13         Sequence 11,           3         4.3         2.3         3         US-10-899-422-13         Sequence 11,           3         4.3         2.3         3         US-10-899-422-13         Sequence 11,           4         4.4         7.76         4         US-10-99-422-13         Sequence 11,           3         4.3         2.3         3         US-10-99-422-13         Sequence 11,           4         4.4         7.76         4         US-10-99-94-43-4	14	733	16.5	155	4	10-106-698-	
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208 4.7 235 3 US-09-894-749-2 Sequence 2, 208 4.7 235 4 US-10-258-3718-20 Sequence 20, 208 4.7 235 4 US-10-258-3718-20 Sequence 20, 208 4.7 235 5 US-10-989-054-2 Sequence 13, 2198.5 4.5 1059 5 US-10-899-422-11 Sequence 13, 2193 4.3 235 5 US-10-899-422-11 Sequence 11, 21, 21, 21, 21, 21, 21, 21, 21, 21,	16	208	4.7	227	٣	US-09-867-550-848	
8 208 4.7 235 4 US-10-258-371B-20 Sequence 20, 208 8.7 235 5 US-10-989-054-2 Sequence 2, 20, 208 8.7 235 5 US-10-989-052-13 Sequence 12, 20, 20, 20, 20, 20, 20, 20, 20, 20, 2	17	208		235	m	US-09-894-749-2	7
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	27		4.3	519	4	-10-428-487-1	14,

Sequence 28, Appl	Sequence 3970, Ap	Sequence 15, Appl	Sequence 107, App	Sequence 107, App	Sequence 152, App	Sequence 1, Appli	Sequence 10101, A	Sequence 385, App	Sequence 4, Appli	Sequence 24, Appl	4	Sequence 1507, Ap	Sequence 812, App	387,	Sequence 26, Appl	Sequence 3, Appli	Sequence 339, App
US-10-258-371B-28	US-10-108-260A-3970	US-10-487-092-15	US-10-087-684-107	US-10-218-779-107	US-10-114-270-152	US-10-113-794A-1	US-11-097-143-10101	US-10-473-127-385	US-09-206-639-4	US-10-258-371B-24	US-10-408-765A-493	US-09-925-300-1507	US-09-925-297-812	US-10-473-127-387	US-10-258-371B-26	US-09-206-639-3	US-09-736-457-339
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189.5	189.5	189.5	184.5	184.5	182	180	175.5	174.5	172.5	172.5	172.5	172.5	168	168	167	166	166
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## ALIGNMENTS

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APPLICANT: Ginsburg, Wendy M.
APPLICANT: Ginsburg, Wendy M.
APPLICANT: Ginsburg, Wendy M.
APPLICANT: Ginsburg, Wendy M.
TILE CANT: Albert
TITLE OF INVENTION: Methods of Diagnosis of Soft Tissue Sarcoma, Compositions of TITLE OF INVENTION: Methods for Screening for Soft Tissue Sarcoma Modulators
TITLE OF INVENTION: Methods for Screening for Soft Tissue Sarcoma Modulators
FILE REFERENCE: 05882.0193 NPUS01
CURRENT APPLICATION NUMBER: 2013.11-26
PRIOR PILING DATE: 2003-11-26
NUMBER OF SEQ ID NOS: 8933
SOFTWARE: Patentin version 3.2
SEQ ID NO 1797
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         61 GLGEPEGRASPDSPLTRWTKSLHSLLGDQDGAYLFRTFLEREKCVDTLDFWFACNGFRQM 120
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Sequence 1797, Application US/10723860
Publication No. US20040253606A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                      Best Local Similarity 89.7
Matches 758; Conservative
                                                                                                                                                                                                                                                                                                                                         TYPE: PRT
ORGANISM: Homo sapiens
US-10-723-860-1797
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Patent No. US20010052137A1

GENERAL INFORMATION:

GENERAL INFORMATION:

APPLICANT: KLEIN,

TITLE OF INVENTION: PAXIN DOMAIN-LIKE POLYPEPTIDE INHIBITORS OF GLYCOGEN

TITLE OF INVENTION: SYNTHASE KINASE 3(BETA) ACTIVITY AND ACTIVATORS OF W

TITLE OF INVENTION: SYNTHASE KINASE 3(BETA) ACTIVITY AND ACTIVATORS OF W

TITLE OF INVENTION: SYNTHASE KINASE 3(BETA) ACTIVITY AND ACTIVATORS OF W

CURRENT PAPLICATION NUMBER: US/09/798, 831

CURRENT PAPLICATION NUMBER: US/09/798, 831

CURRENT PAPLICATION NUMBER: US/09/798, 831

PRIOR APPLICATION NUMBER: US/09/798, 831

PRIOR PELING DATE: 2000-03-01
421 HPLALLPSGSYEEDPOTILDDHLSRYLKTPGCOSPGVGRYSPRSRSPDHHOHHHHQOCH
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                                             VMEENAYQVFLTSDIYLEYVRSGGENTAYMSNGGLGSLKVLCGYLPTLNBEBEWTCADLK
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US-09-798-831-8
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LENGTH: 842
TYPE: PRT
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APPLICANT: Wyeth
APPLICANT: Martinez, Robert
APPLICANT: Martinez, Robert
APPLICANT: Brown, Eugene
APPLICANT: Liu, Wei
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR DIAGNOSING AND TREATING COLON
TITLE OF INVENTION: CANCERS
FILE REFERENCE: AM10927 (131996-002000)
CURRENT FILING DATE: 2003-01-06
PRIOR APPLICATION NUMBER: US Provisional Application 60/438,000
PRIOR PILING DATE: 2003-01-06
NUMBER OF SEQ ID NOS: 54873
SEQ ID NO 116
SEQ ID NO 116
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    EMTPVEPAAFAAELISRLEKLKLELESRHSLEERLQQIREDEEKEGSEQALSSRDGAPVQ
                                                                                                TLLSTGGKLPPVA----ACPLLGGKSFLTKQTTKHVHHYIHHHAVPKTKEBIEAEATQR
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89.7%; Pred. No. 1.1e-276;
ive 31; Mismatches 49;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; Sequence 116, Application US/10751736
; Publication No. US20040265230A1
; GENERAL INFORMATION:
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Matches 758
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                                                                                                                                                                                                           EPEGRAS PDS PLTRWTKS LHS LLGDQDGAYLFRTFLEREKCVDTLDFWFACNGFRQMNLK 123
                                                                                                                                                                                                                                                                                      DIKT---LRVAKAIYKRYI-ENNSVVSKQLKPATKTYIRDGIKKQQIGSVMFDQAQTEIQ 179
                                                                                                                                                                                                                                                                                                                                                                  236
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     EGDDGDVSSGPSV-ISHK---LPSGPPMHHFNSRYSETGCVGMQIRDAHEENPESILDEH 476
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               : ||:||||||||| ||:|||||| ||:||||| || VQRVMKTPGCQSPGTGRHSPKXBSGNL 536
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          443 LSRVLKTPGCQSPGVGRYSPRSRSPDHHHQ------HHHHQQCHTLLSTGGKL 489
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GASGHSRSVARAHPFTQDPAMPPLTPPNTLAQL-EEACRRLAEVSK-----PQKQRCCVA 715
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                550 YSK-----CKSHPKAPEPLPGEQFCGSRGGTLPKRNAKGTEPGLALSARDGGMSSAAGGP 604
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                                                                                                                                                           AVMEENAYQVFLTSDIYLEYVRSGGENTAYMS--NGGLGSLKVLCGYLPTLNEEEEWTC-
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                                                                                          Gaps
                                                                                          149;
                                                    Length 842;
                                                Query Match 37.2%; Score 1654.5; DB 3; Length Best Local Similarity 42.5%; Pred. No. 2.8e-108; Matches 385; Conservative 125; Mismatches 246; Indels
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; ORGANISM: Xenopus
US-09-798-831-8
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RESULT

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DIAGNOSING AND TREATING AUTOIMMUNE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MFDQAQTELQAVMEENAYQVFLTSDIYLEYVRSGGENTAYMS--NGGLGSLKVLCGYLPT
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                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                             Indels 155;
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                                                                                                                                                                                                                                                                                                                                           DB 4;
                                                                                                                                                                                                                                                                                                                                         36.4%; Score 1619.5; DB 4; 41.9%; Pred. No. 8.9e-106; ative 127; Mismatches 251;
                                                                                                             FOR
Sequence 35, Application US/10786720
Sequence 35, Application US/10786720
Publication No. US20040191818A1
GENERAL INFORMATION:
APPLICANT: Weth
APPLICANT: O'Toole, Margot
TITLE OF INVENTION: DISEASES
TITLE OF INVENTION: DISEASES
FILE REPERRINCE: 031896-023000 (AM101331L)
CURRENT FILING DATE: 2004-02-26
NUMBER OF SEQ ID NOS: 21135
SOFTWARE: Patentin version 3.2
LENGTH: 862
                                                                                                                                                                                                                                                                                                                                                                             Conservative 127;
                                                                                                                                                                                                                                                                                    ) ORGANISM: Homo sapiens
US-10-786-720-35
                                                                                                                                                                                                                                                                                                                                                           Similarity
                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Simi
Matches 385;
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                                                                                                                                                                                                                                                                                                                                      INEEEEWTC-----ADLKCKLSPTVVGLSSKTLRATASVRSTETAENGFRSFKR 276
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    347
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     348 -EPVNPYYVNAGYALAPATSANDSEQOSLSSDA---DTLSLTDSSVDGIPPYRI--RKQH 401
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PPVAACPLLGGKSFLTKQTTK------HVHHHYIHHHAVPKTKEEIEAEATQRVRC 539
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      540 LCPGGTDYYCYSK-----CKSHPKAPEPLPGEQFCGSRGGTLPKRNAKGTEPGLALSARD 594
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GGMSSAAGGPQLPG--EEGDRSQDVWQWMLESERQ---SKSKPHSAQSIRKSYPLESARA 649
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 HALQASELVVTYFFCGEEIPYRRMIKAQSLTLGHFKEQLSKKGNYRYYFKKASDEFACGA 813
                                                                                                                                                                                                                                                                                                                                                                                   CNGFROMNLKDT---KTLRVAKAIYKRYI-ENNSVVSKOLKPATKTYIRDGIKKQQIGSV
                                                                                                                                                                                                                                                                                                                                                                                                                                                  170 MFDQAQTEIQAVMEENAYQVFLTSDIYLEYVRSGGENTAYMS--NGGLGSLKVLCGYLPT
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                                                                                                                                                                                                                                                                                                                   55 ARRNEDGLG-EPEGRASPDSPLTRWTKSLHSLLGDQDGAYLFRTFLEREKCVDTLDFWFA
                                                                                                                                                                                                                 172;
                                                Wrapper or
                                                                                                                                                                                 Length 912;
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                                                                                                                                                                                                                 Indels
                                                                                                                                                                                                                                                12 DPSSSFREDAPRPPVPGEEGETPPCQPSVGKVQSTKPMPVS----
PRIOR FILING DATE: 2001-05-31
PRIOR APPLICATION VMBER: USSN 60/287,424
PRIOR FILING DATE: 2001-04-30
Remaining Prior Application data removed - See File
NUMBER OF SEQ ID NOS: 768
SEQ ID NO 270
LENGTH: 912
                                                                                                                                                                                 Query Match 36.4%; Score 1617; DB 4;
Best Local Similarity 41.4%; Pred. No. 1.5e-105;
Matches 384; Conservative 125; Mismatches 246;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               -- POKORCCVASOORDRNHSAAGOAGASPFANP-
                                                                                                                 TYPE: PRT
CORGANISM: Homo sapiens
US-10-092-900A-270
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                                                    762
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                                                                                                                                      HPWAGPQLRTSVQPSHLFIQDPTWPPHPAPNPLTQLEEARRRLEEEEKRASRAPSKQRYV 731
                                                                                   784
                                                    714 VASQQRDRNHSAAGQAGASPFANP-----SLAPEDHKEPKKLASVHALQASELV
                                                                           VTYFFCGEE I PYRRMLKAQSLTLGHFKEQLSKKGNYRYYFKKASDEFACGAVFBE I WDDE
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PRIOR APPLICATION NUMBER: USSN 60/274,322

PRIOR FILING DATE: 2001-03-08

PRIOR FILING DATE: 2001-04-13

PRIOR PILING DATE: 2001-04-13

PRIOR PILING DATE: 2001-04-13

PRIOR PILING DATE: 2001-04-13

PRIOR PILING DATE: 2001-03-08

PRIOR FILING DATE: 2001-03-07

PRIOR PRIOR PRIOR DATE: 2001-03-07

PRIOR PLING DATE: 2001-03-07

PRIOR PLING DATE: 2001-07-10
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CURRENT FILING DATE: 2002-03-07
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APPLICATION NUMBER: USSN 60/294,899
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Shenoy, Suresh G.
Taupier Jr., Raymond J
Pena, Carol E.A.
Li, Li
                                                                                                                                                                                       TVLPMYEGRILGKVERID 840
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Tchernev,, Velizar T.
Fernandes, Elma R.
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Spaderna, Steven K.
Catterton, Blina
Leite, Mario W.
Zhong, Haihong
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Patturajan, Meera
Gangolli, Esha A.
Vernet, Corine A.M.
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Malyankar, Uriel M.
Gerlach, Valerie
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Gusev, Vladimir Y.
Ji, Weizhen
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ller, Charles E.
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Lepley, Denise M.
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US-10-092-900A-270
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FOR DIAGNOSING AND TREATING AUTOIMMUNE
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                                                                                                                                                                                                                                                  DB 4;
                                                                                                                                                                                                                                               Query Match 36.3%; Score 1612.5; DB 4; Best Local Similarity .42.1%; Pred. No. 2.6e-105; Matches 380; Conservative 124; Mismatches 239;
APPLICANT: Liu, Wei
TITE OF INVENTION: COMPOSITIONS AND METHODS FOR
TITLE OF INVENTION: DISEASES
FILE REFERENCE: 031896-023000 (AMI01331L)
CURRENT APPLICATION NUMBER: US/10/786,720
CURRENT FILING DATE: 2004-02-26
NUMBER OF SEQ ID NOS: 21135
SOFTWARE: PALENTIN VETSION 3.2
SEQ ID NO 36
LENGTH: 826
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US-10-786-720-36
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; OTHER INFORMATION: Xaa equals any of the twenty naturally occurring L-amino acids
US-10-264-049-2846
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    INFORMATION: Xaa equals any of the twenty naturally occurring L-amino acids
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   SAQPCDSIVVAYYFCGEPIPYRTLVRGRAVTLGQFKELLTKKGSYRYYFKKVSDEFDCGV
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          36.3%; Score 1615.5; DB 4; Length 347; 86.5%; Pred. No. 4.7e-106; ive 13; Mismatches 33; Indels 1;
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                                                                                                                                                                                                      APPLICANT: Birse et al.
TITLE OF INVENTION: Nucleic Acids, Proteins, and
FILE REFERENCE: PA139P1
CURRENT APPLICATION NUMBER: US/10/264,049
CURRENT FILING DATE: 2002-10-04
FRIOR APPLICATION NUMBER: PCT/US01/18569
FRIOR FILING DATE: 2001-06-07
FRIOR APPLICATION NUMBER: US 60/209,467
FRIOR FILING DATE: 2000-06-07
NUMBER OF SEQ ID NOS: 4360
SOFTWARE: PatentIn Ver. 3.1
SEQ ID NOS: 2846
                                    840
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886 VFEEVREDEAVLPVFEEKIIGKVEKVD 912
                                    814 VFEEIWDDETVLPMYEGRILGKVERID
                                                                                                                                                        Sequence 2846, Application US/10264049 Publication No. US20040005579A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; Sequence 36, Application US/10786720; Publication No. US20040191818A1; GENERAL INFORMATION: APPLICANT: Wyeth; APPLICANT: O'Toole, Margot
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Best Local Similarity 86.59
Matches 300; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ORGANISM: Homo sapiens
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OTHER INFO
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TITLE OF INVENTION: Reagents and Method for Modulating DKK-Mediated Interactions FILE REFERENCE: 032796-143
                                                             PPVAACPLIGGKSFLTKQTTK------HVHHYIHHHAVPKTKEEIEAEATQRVRC 539
                                                                                                                                                           650 APGERVSRHHLLGASGHSRSVARAHPFTQDPAMPPLTPPNTLAQLEEACRRLAEVSK--- 706
                                                                                                                                                                                                                                                                                                             ----LSLBHPWAGPQLRTSVQPSHLFIQDPTMPPHPAPNPLTQLBEARRKLEBBEKRAS 760
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                                                                                    LCPGGTDYYCYSK-----CKSHPKAPEPLPGEQFCGSRGGTLPKRNAKGTEPGLALSARD
                                                                                                                                                                                                          595 GGMSSAAGGPQLPG--EEGDRSQDVWQWMLESERQ---SKSKPHSAQSIRKSYPLESARA
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              12 DPSSSFREDAPRPPVPGEEGETPPCOPSVGKVQSTKPMPVS
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Best Local Similarity 41.2%; Pred. No. 1e-104;
Matches 382; Conservative 125; Mismatches 248;
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CURRENT FILING DATE: 2002-08-02
PRIOR APPLICATION NUMBER: PCT/US02/15982
PRIOR FILING DATE: 2002-06-17
PRIOR FILING DATE: 2001-05-17
PRIOR PILING DATE: 2001-05-17
PRIOR APPLICATION NUMBER: US 60/353,058
PRIOR PILING DATE: 2002-02-01
PRIOR PILING DATE: 2002-03-04
PRIOR PILING DATE: 2002-03-04
PRIOR FILING DATE: 2002-03-04
NUMBER OF SEQ ID NOS: 216
SOFTWARE: FASISEQ for Windows Version 4.0
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874 VFEEVREDEAVLPVFEEKIIGKVEKVD 900
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Robinson, John
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LENGTH: 900
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TITLE OF INVENTION: THE HIGH BONE MASS GENE OF 11q13.3
FILE REFERENCE: 032796-021
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Best Local Similarity 41.2%; Pred. No. 1e-104;
Matches 382; Conservative 125; Mismatches 248;
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CURRENT PAPLICATION NUMBER: US/10/374,979
CURRENT FILING DATE: 2003-03-04
PRIOR PELING DATE: 2000-04-05
PRIOR FILING DATE: 2000-04-05
PRIOR FILING DATE: 2000-04-05
PRIOR PLING DATE: 2000-04-05
PRIOR PLING DATE: 1900-01-03
PRIOR PLING DATE: 1998-01-13
PRIOR PELING DATE: 1998-01-13
PRIOR FILING DATE: 1998-01-13
PRIOR FILING DATE: 1998-01-13
PRIOR FILING DATE: 1998-01-13
PRIOR FILING DATE: 1998-10-23
SEQ ID NO 91
                                                                                                                                                                                                               ; Sequence 91, Application US/10374979; Publication No. US20030219793A1; GENERAL INFORMATION:
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PRIOR APPLICATION NUMBER: US 60/290,071
PRIOR FILING DATE: 2001-05-11
PRIOR PELING DATE: 2001-05-17
PRIOR FILING DATE: 2001-05-17
PRIOR FILING DATE: 2002-02-01
PRIOR PILING DATE: 2002-03-04
PRIOR APPLICATION NUMBER: US 60/351,058
PRIOR PILING DATE: 2002-03-04
NUMBER: PERSER OF SEQ ID NOS: 812
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 670
LENGTH: 900
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US-10-477-238A-670
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  CNGFROMNLKDT --- KTLRVAKAIYKRYI - ENNSVVSKQLKPAIKTYIRDGIKKQQIGSV 169
                                                                                                                                  LNEEEEWTC-----ADLKCKLSPTVVGLSSKTLRATASVRSTETAENGFRSFKR 276
                                                                                                                                                                                SDPVNPYHVGSGYVFAPATSANDSE----LSSDALTDDSMSMTDSSVDGVPPYRMGSKKQL 333
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                                                   CTGFRKLEPCDSNEEKRLKLARAIYRKYILDNNGIVSRQTKPATKSFIKGCIMKQLIDPA
                                                                                                 218 MFDQAQTEIQATMEENTYPSFLKSDIYLEYTRTGSESPKVCSDQSSGSTGKGISGYLPT
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APPLICANT: Yaworsky, Paul
APPLICANT: Yaworsky, Paul
APPLICANT: Bex, Frederick J. III
APPLICANT: Bodine, Peter Van Nest
TITLE OF INVENTION: Transgenic Animal Model of Bone Mass Modulation
FILE REPERENCE: 032796-212
CURRENT APPLICATION UNBER: US/10/477,238A
CURRENT FILING DATE: 2003-11-10
                                                                                                                                                                                                                                                                                                                                        RLQQIREDEEKEGSEQALSSRDGAPVQ-----HPLALLPS--
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VFEEVREDEAVLPVFEEKIIGKVEKVD 900
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AHEENPESILDEHVQRVLRTTGRQSPG-----PGHRSPDSGHV-------ARM
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                                   Indels 172;
Length 900;
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Db 390 RREMQESAQVNGRVPLPHIPRTYRVPKEVR-VEDQKFAEELIHRLEAVORTREAEEKLEE 448  Qy 394 RLQQIREDEEKEGSEQALSSRDGAPVQHPLALLPSG 429        ::	541 PVALGGAASGHGKHYPKSGAKLDAAGL 540 LCPGGTDYYCYSKCKSHPKAPE 549 SFAWGLEPHSHGARSRGYSESVGAAPN 595 GGMSSAAGGOLPGEEGDRSQDVWQ 556EVPGASEDAEKNQKIMQ 650 APGERVSRHHLLGASGHSRSVARAHPF 706LŠLEHPWAGPQLRTSVQPSHLF	Qy         707PQKQRCCVASQQRDRNHSAAGQAGASPFANPSLAPEDHKEPKKLASV 753           Db         761 RAPSKQRYVQEVMRRGRACVRPACAPULHVVPAVSDMELSETETRSQRKVGGG 813           Qy         754 HALQASELVVTYPFCGEBIPYRRMLKAQSLTLCHFKEQLSKKGNYRYYPKKASDEFACGA 813           Qy         754 HALQASELVVTYPFCGEBIPYRRMLKAQSLTLCHFKEQLSKKGNYRYYPKKASDEFACGA 813           Db         814 SAQPCDSIVVAYYFCGEPIPYRTLVRGAAVTLCGFKELLTKKGSYRYYPKKVSDEFDCGV 873           QY         814 VPEEIWDDETVLPMYEGRILGKVERID 840           Db         874 VPEEVREDEAVLPVFEEKIIGKVERVD 900           RESULT 12 US-10-477-173-670         5 Sequence 670, Application US/10477173	Publication No. US20050070699A1   GENERAL INFORMATION:   APPLICANT: Genome Therapeutics Corporation and   APPLICANT: Allen, Kristina M.     APPLICANT: Allen, Kristina M.     APPLICANT: Morales, Arturo J.     APPLICANT: Morales, Arturo J.     APPLICANT: Anisowicz, Anthony     APPLICANT: Anisowicz, Anthony     APPLICANT: Liu, Wei     TILE REFERENCE: 032796-135     CURRENT APPLICATION NUMBER: US/10/477,173     CURRENT APPLICATION NUMBER: US 60/290,071     PRIOR FILING DATE: 2003-11-10     PRIOR FILING DATE: 2001-05-11     PRIOR FILING DATE: 2001-05-17     PRIOR FILING DATE: 2001-05-17     PRIOR PRILICATION NUMBER: US 60/391,311	prior Filing DATE: 2002-02-01  prior APPLICATION NUMBER: US 60/361,293  prior APPLICATION NUMBER: US 60/361,293  prior Filing DATE: 2002-03-04  copymark: FastSEQ for Windows Version 4.0  seq ID NO 670  IENGTH: 900  TYPE: PRT  coranish: Homo sapiens  US-10-477-173-670  Query Match  Best Local Similarity 41.2%; Pred. No. 1e-104;
Db 761 RAPSKQRYVQEVMRRGRACVRPACAPVLHVVPAVSDMELSETETRSQRKVGGG 813  Qy 754 HALQASELVYTYFFCGEEIPYRRMLKAQSLTLGHFKEQLSKKGNYRYYFKKASDEFACGA 813	RESULT 11 US-10-680-287A-670 ; Sequence 670, Application US/10680287A ; Publication No. US20040244069A1 ; GENERAL INFORMATION: ; APPLICANT: Babij, Philip ; APPLICANT: Babij, Frederick J. III ; APPLICANT: Bear, Frederick J. III ; APPLICANT: Bear, Strederick J. III ; APPLICANT: APPLICANT: APPLICANT: ; TILLE OF INVENTION: Transgenic Animal Model of Bone Mass Modulation ; FILE REPERENCE: 032796-179 ; CURRENT APPLICATION NUMBER: US/10/680,287A ; CURRENT FILING DATE: 2003-10-08	PRIOR APPLICATION NUMBER: PCT/US02/14876  PRIOR FILING DATE: 2002-05-13  PRIOR PILING DATE: 2001-05-11  PRIOR PELING DATE: 2001-05-11  PRIOR PELING DATE: 2001-05-17  PRIOR PELING DATE: 2001-05-17  PRIOR PELING DATE: 2001-05-17  PRIOR PELING DATE: 2002-02-01  PRIOR PELING DATE: 2002-02-01  PRIOR PELING DATE: 2002-03-04  NUMBER OF SEQ ID NOS: 812  SOFTWARE: FASTSEQ for Windows Version 4.0  SOFTWARE: PRIOR OF OR WINDOWS O	Query Match   36.1%;   Score 1605;   DB 5;   Length 900;     Best Local Similarity   41.2%;   Pred. No. 1e-104;     Matches   382;   Conservative   125;   Mismatches   248;   Indels   172;   Gaps   29;     Qy	218 LNEEEEWTCADLKCKLSPTVGSESPKVCSDQSSGSGTGKGISGYLPT 228 LNEEEEWTCADLKCKLSPTVGLSSKTLRATASVRSTETAENGFRSFKR 228 LNEEEEWKCDQDMDEDDGRDAAPPGRL-PQKLLLETAAPRVSSRRYSEGREFRYGSWR

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                   12 DPSSSFREDAPRPPVPGEEGETPPCQPSVGKVQSTKPMPVS
Mismatches 248;
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VFBEVREDEAVLPVFEEKIIGKVEKVD 900
Conservative 125;
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; Sequence 34, Application US/10786720; Publication No. US20040191818A1; GENERAL INFORMATION:

APPLICANT: Wyeth

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Cancer Associated Polynucleotides and Polypeptic
                                FOR DIAGNOSING AND TREATING AUTOIMMUNE
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19.3%; Score 860; DB 4; I
Best Local Similarity 50.5%; Pred. No. 2.9e-52;
Matches 192; Conservative 50; Mismatches 82;
APPLICANT: Liu, Wei TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR FILE OF INVENTION: DISEASES.
FILE REPERENCE: 031896-023000 (AMI01331L); CURRENT APPLICATION NUMBER: US/10/786,720; CURRENT FILING DATE: 2004-02-26; NUMBER: PALENTIN VERSION 3.2; SOFTWARE: PALENTIN VERSION 3.2; SEQ ID NOS: 21135; SEQ ID NOS: 21000 3.4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: Colon and Colon Cancer A TITLE OF INVENTION: Colon and Colon Cancer A FILE REFERENCE: PA005P1
CURRENT APPLICATION NUMBER: US/10/106,698
CURRENT FILING DATE: 2002-03-27
PRIOR APPLICATION NUMBER: PCT/US00/26524
PRIOR PILING DATE: 1999-09-29
PRIOR PILING DATE: 1999-09-29
PRIOR FILING DATE: 1999-09-29
PRIOR FILING DATE: 1999-11-03
NUMBER OF SEQ ID NOS: 8564
SEQ ID NO 5828
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CRGANISM: Homo sapiens
US-10-786-720-34
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ORGANISM: Homo
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    equals any of the naturally occurring L-amino acids
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EXPRESSION OF 10,000 OR MORE
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MANE/KEY: MISC FEATURE

JOCATION: (7) CARTON: Xaa equals any of the naturally occurring L-amino and the naturally occurring L-amino and other information: Xaa equals any of the naturally occurring L-amino and CATION: (12)

JOCATION: (12)

OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino and CATION: (48)

LOCATION: (48)

COHER INFORMATION: Xaa equals any of the naturally occurring L-amino and CATION: (48)

COHER INFORMATION: Xaa equals any of the naturally occurring L-amino and US-10-106-698-5828
                                                                                                                                                                                                                                                                         Gaps
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                                                                                                                                                                                                                                 Length 155,
                                                                                                                                                                                                                                                                         11; Indels
                                                                                                                                                                                                                                 Score 733; DB 4;
Pred. No. 6.4e-44;
3; Mismatches 11;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FFACGAVFEEIWEDETVLPMYEGRILGKVERID 155
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE REFERENCE: CLOOO728

CURRENT APPLICATION NUMBER: US/11/097,143

CURRENT PILING DATE: 2005-04-04

PRIOR PLICATION NUMBER: 05/11/097,143

PRIOR FILING DATE: 1999-10-05

PRIOR FILING DATE: 1999-10-05

PRIOR FILING DATE: 1999-10-19

PRIOR FILING DATE: 1999-10-19

PRIOR FILING DATE: 1999-10-19

PRIOR FILING DATE: 1999-11-12

PRIOR FILING DATE: 1999-11-12

PRIOR FILING DATE: 1999-11-12

PRIOR FILING DATE: 1999-11-2

PRIOR FILING DATE: 1999-11-2

PRIOR FILING DATE: 2000-01-12

PRIOR FILING DATE: 2000-01-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: DETECTION KIT, SUCH AS TITLE OF INVENTION: ARRAYS, FOR DETECTING TITLE OF INVENTION: DROSOPHILA GENES.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQ ID NOS: 43008
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 3015
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 3015, Application US/11097143
Publication No. US20050208558A1
GENERAL INFORMATION:
APPLICANT: Venter, J. Craig
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PSGIRKHDDNECSGPRPPVPGEE
                                                                                                                                                                                                                                 Cuery Match 16.5%;
Best Local Similarity 90.8%;
Matches 139; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; ORGANISM: DROSOPHILA US-11-097-143-3015
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US-11-097-143-3015
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Best Local S:
Matches 212,
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                                                                                     KTLRVAKAIYKRYIENNSVVSKQLKPATKTYIRDGIKKQ---QIGSVMFDQAQTEIQAVM 182
                                                                                                                                                                               183 EENAYQVFLTSDIYLEYVR-----SG--GENTAYMSNGGLGSLKVLCGYLPTLN 229
                                                                                                                                                                                                                             214
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               335 EIRP-PGAHGYVYNPSTINTSYVPNSRVDSERASVSSGGRIDSDIMSISSCSMDGRPYIQ 393
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     619 QWMLESERQSKSKP-HSAQSIRKSYPLESARAAPG-----ERVSRHHLLGASGHSRSVA 671
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RASPDSPLTRWTKSLHSLLGDQDGAYLFRTFLEREKCV--DTLDFWFACNGFRQMNLKDT 125
                                                                                                                                                                                                                                                                                                                                                                                                          275 İLPTLHEDSVLSLCDDFEKVQMQEGGSLGSGSVGAGARAPDYPIRLTRDLLIATQKRRL 334
                                                                                                                         279 PVNPYHVGSGYVFAPAT-----SANDSE---LSSDALTD-DSMSMTDSSVDGVPPYR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    551 ----HSMKHSKSMPDHSSCSRKLTNKWPSMNTDSGISMFSADTVTKYK--DASSRSGS--
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                            ----WICADLKCKLSPTVVGLSSKTLRATASVRSTE
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ne : 295.821 secs
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Sequence 208, App
                                                                         April 20, 2006, 16:00:23 ; Search time 46.5435 Seconds (without alignments) 794.148 Million cell updates/sec
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                                                                                                                                                         1 MSSAVLVTLLPDPSSSFRED.....DETVLPMYEGRILGKVERID
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1: /SIDS5/ptodata/2/pubpaa/USO8 NEW PUB.pep:*
2: /SIDS5/ptodata/2/pubpaa/USO6.NEW PUB.pep:*
3: /SIDS5/ptodata/2/pubpaa/USO7 NEW PUB.pep:*
4: /SIDS5/ptodata/2/pubpaa/PCT_NEW PUB.pep:*
5: /SIDS5/ptodata/2/pubpaa/USO9 NEW PUB.pep:*
6: /SIDS5/ptodata/2/pubpaa/US10_NEW PUB.pep:*
7: /SIDS5/ptodata/2/pubpaa/US10_NEW PUB.pep:*
8: /SIDS5/ptodata/2/pubpaa/US10_NEW_PUB.pep:*
GenCore version 5.1.7
Copyright (c) 1993 - 2006 Biocceleration Ltd.
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US-11-169-041-234

US-11-132-687-2

US-11-132-687-2

US-11-1096-588A-29371

US-11-096-588A-29370

US-11-096-588A-29370

US-11-096-588A-29370

US-11-096-588A-29370

US-11-096-588A-29370

US-11-096-588A-29370

US-10-310-773-907

US-10-310-773-907

US-10-310-773-907

US-11-098-686-10232

US-11-098-686-10232

US-11-204-639-11

US-11-204-639-11

US-11-295-561-1004

US-10-995-561-1001

US-10-995-561-998

US-10-995-561-998
                                                                                                                                                                                                                                                  hits satisfying chosen parameters:
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Maximum Match 100%
Listing first 45 summaries
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seq length: 200000000
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Match Length DB
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Perfect score:
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APPLICANT: Briescl-Myers Squibb Company
TITLE OF INVENTION: IDENTIFICATION OF POLYNUCLECTIDES AND POLYPEPTIDE FOR PREDICTING
TITLE OF INVENTION: ACTIVITY OF COMPOUNDS THAT INTERACT WITH PROTEIN TYROSINE KINAS
TITLE OF INVENTION: AND/OR PROTEIN TYROSINE KINASE PATHWAYS
FILE REPERENCE: D0185 PCT
CURRENT APPLICATION NUMBER: US/10/501,035
CURRENT APPLICATION NUMBER: US 60/350,061
PRIOR FILING DATE: 2004-07-09
PRIOR FILING DATE: 2002-01-18
NUMBER OF SEQ ID NOS: 795
SOFTWARE: Patentin version 3.2
 305, App
306, App
306, App
434, App
304, App
194, App
194, App
290, App
290, App
28, App
28, App
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                                                                                                               23491, A
3151, Ap
10104, A
184, App
52, Appl
52, Appl
52, Appl
52, Appl
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LNEEEEWTC-----ADLKCKLSPTVVGLSSKTLRATASVRSTETAENGFRSFKR 276
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       54
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       12 DPSSSFREDAPRPPVPGEEGETPPCQPSVGKVQSTKPMPVS-----SN
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US-11-124-368A-305
US-11-124-367A-433
US-11-124-367A-433
US-11-124-368A-306
US-11-124-368A-304
US-11-124-368A-304
US-11-124-368A-291
US-11-124-368A-291
US-11-124-368A-291
US-11-135-855-29
US-11-135-855-29
US-11-135-855-29
US-11-135-855-29
US-11-096-568A-23491
US-11-096-568A-23491
US-11-096-568A-2491
US-11-096-268A-2491
US-11-096-268A-2491
US-11-096-268A-2491
US-10-242-586-52
US-10-242-586-52
US-10-242-384-52
US-10-242-384-52
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 36.1%; Score lous; up 0; Best Local Similarity 41.2%; Pred. No. 2.2e-107; Matches 382; Conservative 125; Mismatches 248;
                                                                                                                                                                                                             ALIGNMENTS
                                                                                                                                                                                                                                                          Sequence 215, Application US/10501035
Publication No. US20060046249A1
GENERAL INFORMATION:
  ; ORGANISM: Homo sapiens
US-10-501-035-215
  US-10-501-035-215
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10232, A

Sequence Sequence Sequence Sequence

Sequence

Sequence Sequence

Sequence

136.5 136.5 136.5 136.5 129.5 128 128 128 128

Sequence Sequence Sequence

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APPLICANT: Habener, Joset
APPLICANT: Abraham, Elizabeth
APPLICANT: Abraham, Elizabeth
APPLICANT: Abraham, Elizabeth
APPLICANT: Abraham, Elizabeth
APPLICANT: Vallejo, Mario
TITLE OF INVENTION: BY ADMINISTERING A PSEUDO-ISLET LIKE AGGREGATE DIFFERENTIATED
TITLE OF INVENTION: BY ADMINISTERING A PSEUDO-ISLET LIKE AGGREGATE DIFFERENTIATED
TITLE OF INVENTION: A NESTIN-POSITIVE PANCREATIC STEM CELL.
FILE REFERENCE: 3284/1233
CURRENT APPLICATION NUMBER: US/10/984,645
CURRENT FILING DATE: 2004-11-09
PRIOR PILING DATE: 1999-12-06
PRIOR PILING DATE: 1999-12-06
PRIOR PILING DATE: 1000-06-28
PRIOR PILING DATE: 1000-06-28
PRIOR PILING DATE: 2000-10-06
PRIOR PILING DATE: 2000-10-06
PRIOR FILING DATE: 2000-10-06
PRIOR FILING DATE: 2000-10-06
PRIOR FILING DATE: 2000-10-06
                                                                                                                                                                                                                                                                       APPLICANT: Bristol-Myers Squibb Company
TITLE OF INVENTION: IDENTIFICATION OF POLYNUCLECTIDES FOR PREDICTING ACTIVITY OF
TITLE OF INVENTION: COMPOUNDS THAT INTERACT WITH AND/OR MODULATE PROTEIN TYROSINE
TITLE OF INVENTION: KINASES AND/OR PROTEIN TYROSINE KINASE PATHWAYS IN LUNG CANCER
TITLE OF INVENTION: CELLS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     86 IGDQDGAYLFRIFLEREKCVDTLDFWFACNGFRQMNLKDTKTLRVAKAIYKRYIENNSVV 145
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                -----WSEAFDEL 87
LGDQDGAYLFRTFLEREKCVDTLDFWFACNGFRQMNLKDTKTLRVAKAIYKRYIENNSVV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           30 EGETPPCOPSVGKVQS----TKPMPVSSNARRNEDGLGEPEGRASPDSPLTRWTKSLHSL
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                                                                               146 SKOLKPATKTYIRDGIKKQQIGSVMFDQAQTEIQAVMEENAYQVFLTSDIY 196
                                                                                                    Length 211;
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26.9%; Pred. No. 3.9
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                                                                                                                                                                                                                                                                                                                                                                              FILE REFERENCE: 10001 NP
CURRENT APPLICATION NUMBER: US/11/169,041
CURRENT FILING DATE: 2005-06-28
PRIOR APPLICATION NUMBER: 60/584,405
PRIOR FILING DATE: 2004-06-30
                                                                                                                                                                                  RESULT 3
34
Sequence 234, Application US/11169041
Sequence 234, Application US/11169041
Publication No. US20060019284A1
GENERAL INPORMATION:
GENERAL INPORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 2, Application US/10984645; Publication No. US20050244386A1; GENERAL INFORMATION: APPLICANT: Habener, Joel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQ ID NOS: 527
SOFTWARE: PatentIn version 3.2
SEQ ID NO 234
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US-10-984-645-2
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TITLE OF INVENTION: IDENTIFICATION OF POLYNUCLEOTIDES AND POLYPEPTIDE FOR PREDICTING
TITLE OF INVENTION: ACTIVITY OF COMPOUNDS THAT INTERACT WITH PROTEIN TYROSINE KINASE
TILLE OF INVENTION: AND/OR PROTEIN TYROSINE KINASE PATHWAYS
FILE REFERENCE: D0185 PCT
CURRENT APPLICATION NUMBER: US/10/501,035
PRIOR TELING DATE: 2004-07-09
PRIOR FILING DATE: 2002-01-18
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Pred. No. 3.9e-05;
24; Mismatches 72;
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SOFTWARE: PatentIn version 3.2
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US-10-501-035-208
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US-10-501-035-208
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                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                       Query Match 3.4%; Score 151.5; DB 6; Length 1618; Best Local Similarity 19.4%; Pred. No. 0.023; Matches 180; Conservative 108; Mismatches 333; Indels 309;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  170 MFDQAQTEIQAVMEENAYQVF------LTSDIYLEYVRS----
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SOFTWARE: PatentIn version 3.2 SEQ ID NO 2
                                                                                                          ORGANISM: Homo sapiens
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                                                LENGTH: 1618
TYPE: PRT
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                                                                                                                                                                                                                               APPLICANT: Habener, Joel
TITLE OF INVENTION: STEM CELLS AND THEIR USE IN TRANSPLANTATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 3.4%; Score 151.5; DB 7; Best Local Similarity 19.4%; Pred: No. 0.023; Matches 180; Conservative 108; Mismatches 333;
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1181 RGAEEAFPAETLGHTGSDAPSPWPLGSEEA 1210
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                                                                                                                                                                                                                                                                                FILE REFERENCE: 3284/1238
CURRENT APPLICATION NUMBER: US/11/132,687
CURRENT FILING DATE: 2005-05-19
PRIOR APPLICATION NUMBER: US 60/16,082
PRIOR APPLICATION NUMBER: US 60/215,109
PRIOR PILING DATE: 2000-06-28
PRIOR FILING DATE: 2000-06-28
PRIOR FILING DATE: 2000-10-06
PRIOR FILING DATE: 2000-10-06
PRIOR FILING DATE: 2000-110-06
PRIOR FILING DATE: 2000-110-06
                                                                                                                                        Sequence 2, Application US/11132687; Publication No. US20060062769A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: Patentin version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: PRT
ORGANISM: Homo sapiens
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518 Db 2266 VSGSPAPSTSGTSTPRRGRRQLPQTPSTPRPHVSYSPVIRKAGGSGPPQQQQOOQQQQ 2323. 992 Qy 626 RQSKSKP-HSAQSIRKSYPLESARAAPGERVSRHHLLGASGHSRSVARAHPFTQDPAMPP 684	578   Db 2323 QQAVARPGRAATSGPRRYPGPTAEPLAGDRPPTGGHSSGRSPRMER 2368	1028 Oy 685 LIPPNTLAQLEEACR 699	638 Db 2369 RVPGPARSESPRACR 2383	1064 RESULT 7 682 US-11-096-568A-29371 ; Sequence 29371, Application US/11096568A ; Publication No. US20060048240A1	; GENERAL INFORMATION: ; APPLICANT: Alexandrov, Nickolai et al. ; TITLE OF INVENTION: Sequence-Determined DNA Fragments and Corresponding Polypeptide: ; TITLE OF INVENTION: Therby	; FILE REFERENCE: 2750-1592PUS2 ; CURRENT APPLICATION NUMBER: US/11/096,568A ; CURRENT FILING DATE: 2005-04-01 . NUMBER OF SED ID NOS: 34471	; NOVIDEN OF SEQ ID NOS: 51471 ; SEQ ID NO 29371 ; LENGTH: 496	CORE US-1			DD 1.25 DELNITRYVEGIDPDIGIRELEEDSDSSTREMFESSVILMGSTLEGITGETTE CONTROL OF STREET		Db 233 INLAPYENLYVSLSNSEGSTVSPPTTVHSSVLLRVGTSNSSPRLKQLTDT1TGSRSKNLG	2047 . Qy 437TILDDHLSRVLKTPGCQSPGVGRYSPRSRSPDHHHQACHT 481	482 LLSTGGKLPPVAACPLLGGKSFLTKQTTKHVHHY	404 Db 346	4 Db 378	2201 QY SSAAGGPQLPGEEGDRSQDVWQWMLESERQSKSKPHSAQSIRKSYPLESARAAPGERVS	524 Db 413 430 224 Db 417 418 430 224 Db 417 GASGHSRSVARAHPTODPAMPPLTPPNTLAOLEBACRRLAEVSKPQKQ 710	4 Db 431	2265 PERITT A
QY 491 PVAACPLIGGKSFLTKQTTKHVHHYIH	FCGSRGGTLPK	DD :   :       :   :   :   :   :   :   :	579 RNAKGTEPGLALSARDGGMSSAAGGPQLPGEEGDRSQDVWQWMLESERQSKSKPHSAQSI	Db 1029 ASVKGGABGILQDPEGQSQQVGTPGLQAPQG	QY 683 -PPLTPPNTLAQLEEACRRLAEVSK-PQKQRCCVASQQRDRNHSAA	727	DD 1181 KGAEEAFPAETLGHIGSDAFSFWFLGSEEA 1210	RESULT 6 US-11-126-313-33 Sequence 33, Application US/11126313 Publication No. US20050288489A1 GENERAL INFORMATION: A APPLICANT HISCH, OG A TTHE OF THY SYNTYN. VOLTARE-DREPENDENT CALCITM CHANNEL BETA SUBINIT FUNCTIONAL TTHE OF THY SYNTYN. VOLTARE-DREPENDENT CALCITM CHANNEL BETA SUBINIT FUNCTIONAL	ei	ARE: Patentin ve. NO 33 TH: 2505	; TYPE: PRT ; ORGANISM: homo sapiens US-11-126-313-33	3.4%; Score 149; DB 7; Length 2505; hrity 20.8%; Pred. No. 0.065;	I//; Indels 168; Gaps SELSSDALTDDSMSMTDS	1988 FORMEPPSPTQEGGPGNALPSTQLDPGGALMAHESGLKESPSWVTQRAQEMPQKTGTWS	DY 318 SVDCVPFYRMSSKKULG	359 PKEMTPVEPAAPABLISRLEKLKLELESRHSLEERLQQIREDBEK	COSPGVGRYSPRS	2162 RASERSLGRY-TDVDTGLGTDLSMTTQSGDLPSKERDQERG	465 RSPDHHHQHHHQQCHTLLSTGGKLPPVAACPLLGGKSFLTKQTTKHHHYIHHAVPK	TKEEI EAEATQRVRCLCPGGTDYYCYSKCKSHPKAPEPLPGEQFCGSRGGTLPKRNAKGT	DD 2225 DKDRYAQERPDHGRSR-RDQRWSRSFSEGREHMAHRQGSSS

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24;
                                                                                                                                                                                                                                                     313 DELNITKVVFGIDPDTGYREILPLSLSSIK----EMFESVLINQSTLQLTKSLFGETFLF 369
                                                                                                                                                                                                                     310 DSMSMT-----DSSVDGVPPYRMGSKKQLQREMHRSVKANG---QVSLPHFPRTH-- 356
                                                                                                                                                                                                                                                                                                                                                        369 EVLKFPGGITVIPPQSAFP-----LQKFKIVF--NFTLNYSIHQIQINFNTLASQLKNG 420
                                                                                                                                                                                                                                                                                                                                                                                                                                                        421 INLAPYENLYVSLSNSEGSTVSPPTTVHSSVLLRVGTSNSSPRLKQLTDTTTGSRSKNLG 480
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LLSTGGKLPPVAACPLLGGKSFLTKQTTKHVHHYIHHAVPKTKEEIE---AEATQRVR 538
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               481 LNNTIFGKVKOVRLSSFLPNSSDSSTKSPS-PSP----SPHSKHNHHHHHHHHHHH 533
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               539 CLCPGGTDYYCYSKCKSHPKAPEPLPGEQFC--GSRGGTLPKRNAKGTEPGLALSARDGG 596
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                597 MSSAAGGPQLPGEEGDRSQDVWQWMLESERQSKSKPHSAQSIRKSYPLESARAAPGERVS 656
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ----TILDDHLSRVLKTPGCQSPGVGRYSPRSRSPDHHHQHHHQQCHT
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APPLICANT: Stache-Crain, Birgit
APPLICANT: Andarmani, Susan
APPLICANT: Andarmani, Susan
APPLICANT: Tang, Y. Tom
TITLE OF INVENTION: Methods for Diagnosis and Treatment of Preeclampsia
FILE REFERENCE: 821A
CURRENT APPLICATION NUMBER: US/10/821,234
CURRENT PAPLICATION NUMBER: US/60/462,047
PRIOR APPLICATION NUMBER: US/60/462,047
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            -----GAPH--HQLHSPAPISAAKS-----
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                                                                                                                     Query Match 3.3%; Score 148; DB 7; Length 684; Best Local Similarity 20.7%; Pred. No. 0.012; Matches 98; Conservative 42; Mismatches 132; Indels 202;
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21.5%; Pred. No. 0.01;
iive 64; Mismatches 178;
; NAME/KEY: misc_feature
; LOCATION: (1)...(684)
; OTHER INFORMATION: Ceres Seq. ID no. 4814587
US-11-096-568A-29369
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 1039, Application US/10821234
Publication No. US20050255114A1
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SOFTWARE: pt SEQ genes Version 1.0
SEQ ID NO 1039
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Query Match
Best Local Similarity 21.5
Matches 104; Conservative
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                                                                        and Corresponding Polypeptides
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TITLE OF INVENTION: Sequence-Determined DNA Fragments and Corresponding Polypeptides
TITLE OF INVENTION: Therby
FILE REFERENCE: 2750-1592PUS2
CURRENT APPLICATION NUMBER: US/11/096,568A
CURRENT FILING DATE: 2005-04-01
NUMBER OF SEQ ID NOS: 34471
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        / Match 3.3%; Score 148; DB 7; Length 548;
Local Similarity 20.7%; Pred. No. 0.0088;
nes 98; Conservative 42; Mismatches 132; Indels 202;
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                                       APPLICANT: Alexandrov, Nickolai et al.
TITLE OF INVENTION: Sequence-Determined DNA Fragments a TITLE OF INVENTION: Therby
FILE REPERINCE: 2750-1592PUS2
CURRENT APPLICATION NUMBER: US/11/096,568A
CURRENT FILING DATE: 2005-04-01
NUMBER OF SEQ ID NOS: 34471
SEQ ID NO 29370
                                                                                                                                                                                                                                                                                                                                                                                                    no. 4814588
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                                                                                                                                                                                                                                                                                                                                             NAME/KEY: misc_feature
CATTON: (1)..(548)
OTHER INFORMATION: Ceres Seq. ID
US-11-096-568A-29370
                                                                                                                                                                                                                                                                                              ORGANISM: Arabidopsis thaliana
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ORGANISM: Arabidopsis thaliana
FRATURE:
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Best Local S
Matches 98
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--RKHSLIKNDSSPQRCSPAREPQASAPSPPGLHVDPG----RGMGPLPCGSPRLQLSPL 2026
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                                                                                                                                                                                                                 LGGKSFLTKQTTKHVHHYIHHHAVPKTKEEI-----EAEATQRVRC-LCPGGTDYYCY 550
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SPGVGRYSPRSRSPDHHHOHHHHQQCHTLLSTGGKLPPVAACPLLGGKSFLTKQTTKHVH 513
                                                                                                                                                                                                                                                ----SAL------HDFHGHILARTEENIFSHLPLHSQHLTRAPCPLIPIGGIQMVQ
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                                                                                               TLCP--LGRELAPRAHVLSKLEGTTDPGLPRYSPTRRWSPGQAESPPRSAPPGKWAGPGS
                                                                                                                                      -TGGKLPPVAACPL
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                                                                                                                                                                                                                                                                                                                                                                          -LSARDGG-----MSSAAGGPQLPGEEGDRSQDVWQWMLESERQSKSKPHSAQSIRKSYP
                                                                                                                                                                                                                                                                                                                                                                                                                                                       644 LESARAAP-----GERVSRHHLLGASGHSRSVARAHPFTQDP-AMPPLTPPNTLAQL
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APPLICANT: David W. Morris
APPLICANT: Marc Malandro
TITLE OF INVENTION: Novel Compositions and Methods in Cancer
FILE REFERENCE: 529452001300
CURRENT APPLICATION NUMBER: US/10/330,773
CURRENT FILING DATE: 2002-12-27
NUMBER OF SEQ ID NOS: 981
SOFTWARE: FastSEQ for Windows Version 4.0
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20.7%; Pred. No. 0.21;
ive 76; Mismatches 210;
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                                                            TILDDHLSRVL-----KTPGCQSPGVGRYSPRSR-
                                                                                                                                      ---HTLLS----
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Publication No. US20060040262A1
GENERAL INFORMATION:
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Best Local Similarity 20.7'
Matches 120; Conservative
                                                                                                                                      ннониниоос---
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    SVRSTETAENGFRSFKRSDPVN---PYHVGSGYVFAPATSANDSELSSDALTDDSMSMTD 316
                                                                                SSVDGVPPYRMGSKKQLQREMHRSVKANGQVSLPHFPRTHRLPKEMTPVEPAAFAAELIS 376
                                                                                                                    -----PHLERE-----TP-KPRKSGKVKE 263
                                                                                                                                                                                      -----HPLALLPSGSYEEDPQTILDDHLSRVLKTPGCQSPGVGRYSPRSRSPDHHHQ 472
                                                                                                                                                                                                                                                                               324 EGCGTVLAHPRYLQHHIKYO------HLLK--KKYVCPHPSCGRLFRLQKQLLRHAK 372
                                                                                                                                                                                                                                                                                                                      HH--------HHQQCHTLLSTGGKLPPVAACPLLGGKSFLTKQTTK---H 511
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                                                                                                                                                                                                                                                                                                                                                                                                                                470 LAANAGALITSTDILGTNPESLTQPSDG-----QGLPLLPEPLGNSTSGEC-LLLEARGM 523
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                568 FCGSRGGTLPKRNAKGTEPGLALSARDGGMSSAAGGPQLPGEEGDRSQDVWQWMLESERQ 627
                             EWICADL-------KCKLSPTVVGLSSKTLRATASVRSTETAENGFRSFKRSD
                                                                                                                                                           RLEKLKLELESRHSLEERLOQIREDEE---KEG----SEQALSSRDGAPVQ----
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 907, Application US/10330773
; Sequence 907, Application US/10330773
; Publication No. US20060040262A1
; GENERAL INFORMATION:
; APPLICANT: David W. Morris
; APPLICANT: David W. Morris
; APPLICANT: David W. Morris
; TITLE OF INVENTION: Novel Compositions and Methods in Cancer
; FILE REFERENCE: 529452001300
; CURRENT APPLICATION UNMBER: US/10/330,773
; CURRENT FILING DATE: 2002-12-27
; NUMBER F FSEQ ID NOS: 981
; SEQ ID NO 907
; SEQ ID NO 907
; LENGTH: 2392
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Best Local Similarity 20.3%; Pred. No. 0.32;
Matches 145; Conservative 70; Mismatches 274; Indels
                                                                                                                    EEI ---- PFKDDPRDETYK---
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US-10-330-773-907
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Db	OY 404 KEGSEQALSSRDGAPVQHPLALLPSGSYEEDPQTILDDHLSRVLKTP 450	Qy         451 GCQSPGVGRYSРRSRS,	OY 479 CHTLLSTGGKLPPVAACPLLGGKSFLTKQTTKHVHHYIHHAVP 523	Oy 524 KTKEEIBAEATQRVRCLCPGGTDYYCYSKCKSHPKAPEP 562	LPGEOFCGSRGGT	Query Match 3.1%; Score 137; DB 6; Length 748; Best Local Similarity 35.4%; Pred. No. 0.085; Matches 40; Conservative 16; Mismatches 47; Indels 10; Gaps 5;
Db 1377 AGSVDSQGPAQQEPKHACDTCGKNPKFLGTLSR 1409  Qy 514 HHYIHHAVPKTKEEIBAEATQRVRCLCPGGTDYYCYSKCKSHPKAPEPLPGEQFC 569	Qy 570GSRGGTLPKRNAKGTEPGLALSARDGGMSSAAGGPQLPGEEGDRSQDVWQ- 619	Qy 620WMLES-ERQSKSKPHSAQSIRKSYPLESARAAPGERVSRHHLLGAS 664  Db 1517 CNKRFWSLQDLTRHNRSHTGERPYKCQTCERTFTLKHSLVRHQRIHQKARHSKHHG 1572	Qy 665 GHSRSVARAHPFTQDPAMPPLTPPNTIAQLEEACRRIAEVSKPQKQRCCVASQQRDRNHS 724	Qy         725 AAGQAGASPFANPSLAPEDHKEPKKLASV 753           :	Sequence 44, Application US/10469469	Qy 288GYVPAPATSANDSELSSDALTDDSMSMTDSSVDGVPPYRMGSKKQLQREMHRSVKA 343

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              AAGOAGASPFANPS--LAPEDHKEPKKLASVHALQASELVVTYFFCGEEIPY--RRMLKA 780
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                                    195 IYLEYVRSGGENTAYMSNGGL-----GSLKVLCGYLPTLNEEBEWTCADLKCKLSPTVV
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                                                                                    QSLTLGHFKEQLSKKGNYRYYFKKASDEFACGAVFEEIWDDETVLPMYEGRIL 833
                                                                                                     ERVILADFKGVL-QRPSYKFFFKSMDDDF--GVVKEEISDDNAKLPCFNGRVV 107
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TITLE OF INVENTION: Genes and Uses for Plant Improvement
FILE REFERENCE: 38-21(53450)B EP
CURRENT APPLICATION NUMBER: US/11/087,099
CURRENT FILING DATE: 2005-03-22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              3.1%; Score 136.5; DB 7;
18.7%; Pred. No. 0.12;
tive 96; Mismatches 295;
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Sequence 950, Application US/11087099
Publication No. US20060041961A1
GENERAL INFORMATION:
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TYPE: PRT
ORGANISM: Cochliobolus carbonum
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Matches 147; Conservative
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Sequence

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Sequence 3 Sequence 1 Sequence 1 Sequence 1

Sequence Sequence Sequence

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Sequence

Sequence 1, A Sequence 25,

ALIGNMENTS

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58 YKRYI-ENNSVVSKQLKPATKTYIRDGIKKQQIGSVMFDQAQTEIQAVMEENAYQVFLTS 116
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CUDNIKK: US
CIP: 10036
COMPUTER READABLE FORM:
MEDIUM TYPE: Flopyy disk
MEDIUM TYPE: Flopyy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/NS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/890,865A
FILING DATE: 10-JUL-1997
CLASSIPICATION: 435
ATCHASTRIANG: JOHN P
REGISTRATION: WHERE: 28,678
REFERENCE/DOCKET NUMBER: 0575/54249
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-08-890-865A-19
Sequence 19, Application US/08890865A
APPLICANT: Constantini, Franklin
APPLICANT: Zeng, Li
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APPLICANT: Zeng, Li
APPLICANT: Constantini, Franklin
APPLICANT: Constantini, Franklin
APPLICANT: Zeng, Li
STREET: 1185 Avenue of the Americas
127 amino acids
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
   STRANDEDNESS: single.
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STATE: New York
COUNTRY: US
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989.497 Million cell updates/sec
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                                 GenCore version 5.1.7
Copyright (c) 1993 - 2006 Biocceleration Ltd.
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/cgn2_6/ptodata/1/iaa/6_COMB.pep:*
/cgn2_6/ptodata/1/iaa/H_COMB.pep:*
/cgn2_6/ptodata/1/iaa/PcTuS_COMB.pep:*
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US-10-113-794A-1
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                                                                                                                                                 - protein search, using sw model
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                23; Indels
                                                                                                                                                                                                                                                     SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/890,865A FILING DATE: 10-UJUL-1997 CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDIUM TYPE: Floppy disk
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
63.5%; Score 406; DB 2;
Best Local Similarity 61.1%; Pred. No. 5.2e-42;
Matches 77; Conservative 22; Mismatches 23;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
APPLICANT: Conetantini, Franklin
APPLICANT: Zeng, Li
TITLE OF INVENTION: AXIN GENE AND USES THEREOF
NUMBER OF SEQUENCES: 23
CORRESPONDENCE ADDRESS:
ADDRESSEE: Cooper & Dunham LLP
STREET: 1185 Avenue of the Americas
AXIN GENE AND USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADDRESSEE: Cooper & Dunham LLP
STREET: 1185 Avenue of the Americas
CITY: New York
STATE: New York
                                                    ADDRESSEE: Cooper & Dunham LLP
STREET: 1185 Avenue of the Americas
CITY: New York
STATE: New York
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                                                                                                                                                    ZIP: 10036
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-
                                                                                                                                                                                                                                                                                                                                                                                                     28,678
                                                                                                                                                                                                                                                                                                                                                                                  NAME: White, John P
RECIGTRATION NUMBER: 28,678
REFERENCE/DOCKET NUMBER: 05
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212)278-0400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        INFORMATION FOR SEQ ID NO: 10: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 855 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (212)391-0526
NUMBER OF INVENTION: AX.
CORRESPONDENCE ADDRESS:
ADDRESSE: Cooper & I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY: US
ZIP: 10036
COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MOLECULE TYPE: Protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              linear
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                                                                                                                                           COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LENGTH:
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    62 YRKYILDSNGIVSRQTKPATKSFIKDCVWKQQIDPAMFDQAQTEIQSTWEENTYPSFLKS 121.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          58 YKRYI-ENNSVVSKQLKPATKTYIRDGIKKQQIGSVMFDQAQTEIQAVMEENAYQVFLTS 116
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 WIKSLHSLLGDQDGAYLFRTFLEREKCVDTLDFWFACNGFRQMNLKDT---KTLRVAKAI 57
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               4; Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      23; Indels
                                                                                                                                                                                            Sequence 1, Application US/08890865A; Patent No. 6307019; GENERAL INFORMATION:
APPLICANT: Zeng, Li
TITLE OF INVENTION: AXIN GENE AND USES THEREOF; NUMBER OF SEQUENCES: 23; CORRESPONDENCE ADDRESS: 3
ADDRESSEE: Cooper & Dunham LLP STREET: 1185 Avenue of the Americas STREET: New York
COUNTRY: New York
COUNTRY: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SCHWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/890,865A FILING DATE: 10-JUL-1997 CLASSIFTCATION: 435 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 407; DB 2;
Pred. No. 4.7e-42;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity 61.9%; Pred. ...
....hes 78; Conservative 21; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
APPLICANT: Constantini, Franklin
APPLICANT: Zeng, Li
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: White, John P
REGISTRATION NUMBER: 28,678
REFERENCE/DOCKET NUMBER: 0557
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212)278-0400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               992 amino acids
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 992 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MOLECULE TYPE: Protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ZIP: 10036.
COMPUTER READABLE FORM:
                                                               117 DIYLEY 122
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STRANDEDNESS: si
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                                                                                                   122 DIYLEY 127
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LENGTH: 235
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Hodge, Martin R.
APPLICANT: Yowe, David
TITLE OF INVENTION: RGS-Containing Molecules and Uses Thereof
FILE REFERENCE: 5800-19, 035800/174680
CURRENT APPLICATION NUMBER: US/09/244,314
CURRENT FILIG DAFE: 1999-02-04
NUMBER OF SEQ ID NOS: 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 900;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           62.8%; Score 401; DB 2;
62.7%; Pred. No. 2.4e-41;
iive 17; Mismatches 26
                                                                                                                                                                                                  0575/54249
   US/08/890,865A
PPLICALLE
PILLING DATE: 10-JUL-LL-
FILLING DATE: 10-JUL-LL-
CLASSIFICATION: 435
ATTORNEY AGENT INFORMATION:
NAME: White, John P
REGISTRATION NUMBER: 28,678
REFERENCE/DOCKET NUMBER: 0575/
TELECOMMUNICATION INFORMATION:
TELEPAK: (212) 391-0526
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
LENGTH: 900 amino acids
LENGTH: 900 amino acids
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Sequence 2, Application US/09244314

; Patent No. 6274362

; GENERAL INFORMATION:
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Matches 79; Conservative
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SEQ ID NO 2
LENGTH: 235
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Best Local Similarity 34.6
Matches 44; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TOPOLOGY: linear MOLECULE TYPE: Protein
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US-09-244-314-2
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194 KSDIYLD 200
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RESULT

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55 KAIYKRYIENNSVVSKQLKPATKTYIRDGIKKQQIGSVMFDQAQTEIQAVMEENAYQVFL 114
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Parent No. 6830914

GENERAL INFORMATION:
APPLICANT: Hodge, Martin R.
TILLE OF INVENTION: RGS-Containing Molecules and Uses Thereof
FILE REFERENCE: 5800-19, 035800/174680

CURRENT APPLICATION WUBBER: US/09/894,749

CURRENT FILING DATE: 2001-06-27

PRIOR FILING DATE: 1999-02-04

NUMBER OF SEQ ID NOS: 4

SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Indels 15;
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Sequence 2, Application US/09498959

Patent No. 6410240

GRUERAL INFORMATION

APPLICANT: Hodge, Martin R.

APPLICANT: Yowe, David

TITLE OF INVENTION: RGS-Containing Molecules and Uses

TITLE OF INVENTION: Thereof

FILE REFERENCE: 5800-19A

CURRENT APPLICATION NUMBER: US/09/498,959

CURRENT FILING DATE: 12000-02-04

EARLIER APPLICATION NUMBER: 09/244,314

SARLIER PILING DATE: 1999-02-04

NUMBER OF SEQ ID NOS: 12

SOFTWARE: FastSEQ for Windows Version 3.0

SEQ ID NO
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         43;
                                                                                                                                                                                                                                                                                                                                                                                                                                            29.2%; Score 186.5; DB 2
34.6%; Pred. No. 3.6e-15;
tive 25; Mismatches 43.
                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity 34.53
Best A4; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity 34.6%
Matches 44; Conservative
                                                                                                                                                                                                                                                                                                                                                                             ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: PRT
ORGANISM: Homo sapiens
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61 YIENNSVVSKQLKPATKTYIRDGIKKQQIGSVMPDQAQTEIQAVMEENAYQVFLTSDIYL 120
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                 GENERAL INFORMATION:
APPLICANT: Hodge, Martin R.
APPLICANT: Yowe, David
TITLE OF INVENTION: RS.
PILE REFERENCE: 5800-19, 035800/174680
CURRENT APPLICATION NUMBER: US/09/894,749
CURRENT APPLICATION NUMBER: 09/244,314
PRIOR APPLICATION NUMBER: 09/244,314
PRIOR FILING DATE: 1999-02-04
NUMBER OF SEQ ID NOS: 4
SOFTWARE: Patentin Ver: 2.0
SEQ ID NO 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 2, Application US/10113794A

Sequence 2, Application US/10113794A

Patent No. 6919313

GENERAL INFORMATION:

APPLICANT: Flanagan et al.

TITLE OF INVENTION: BEPHEN REGULATION OF G-PROTEIN COUPLED

TITLE OF INVENTION: CHEMOATTRACTION

FILE REFERENCE: 2535/106

CURRENT APPLICATION NUMBER: US/10/113,794A

CURRENT FILING DATE: 2002-04-01

NUMBER OF SEQ ID NOS: 6

SOFTWARE: FastSEQ for Windows Version 4.0

SEQ ID NO 2

LENGTH: 219
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27.0%; Score 172.5; DB 2; Length
Best Local Similarity 32.5%; Pred. No. 6e-13;
Matches 40; Conservative 24; Mismatches 56; Indels
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31.7%; Pred. No. 4.9e
ive 28; Mismatches
                                                                                                                                 Sequence 4, Application US/09894749
Patent No. 6830914
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ORGANISM: Homo sapiens
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Matches 39; Conserv
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121 EYV 123
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ORGANISM: Mus sp.
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                                                                                        Sequence 4, Application US/09244314

Sequence No. 6274362

GENERAL INFORMATION:

APPLICANT: Hodge Martin R.

APPLICANT: Yowe, David

TITLE OF INVENTION: RGS-Containing Molecules and Uses Thereof

FILE REPERENCE: 8800-19, 035800/174680

CURRENT FILING DATE: 1999-02-04

NUMBER OF SEQ ID NOS: 4

SOFTWARE: PatentIn Ver. 2.0

SEQ ID NO 4

LINCTH: 235
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27.8%; Score 177.5; DB 2; Length
Best Local Similarity 31.7%; Pred. No. 4.9e-14;
Matches 39; Conservative 28; Mismatches 53; Indels
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CURRENT APPLICATION NUMBER: US/09/498,959
CURRENT FILING DATE: 2000-02-04
EARLIER APPLICATION NUMBER: 09/244,314
EARLIER FILING DATE: 1999-02-04
NUMBER OF SEQ ID NOS: 12
SOFTWARE: FASESEQ for Windows Version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match. 27.8%; Score 177.5; DB 2; Best Local Similarity 31.7%; Pred. No. 4.9e-14; Matches 39; Conservative 28; Mismatches 53;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-09-498-959-4
; Sequence 4, Application US/09498959
; Patent No. 6410240
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
APPLICANT: Hodge, Martin R.
APPLICANT: Yowe, David
  194 KSDIYLD 200
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GRANISM: Mus sp.
US-09-244-314-4
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ORGANISM: Mus sp.
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LENGTH: 235
                                                                                 US-09-244-314-4
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61 YIENNSVVSKQLKPATKTYIRDGIKKQQIGSVMFDQAQTEIQAVMEENAYQVFLTSDIYL 120
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                                                                                                                                                                                                                                                                                                                                                                                                                        1 WIKSLHSLLGDQDGAYLFRIFLEREKCVDILDFWFACNGFRQMNLKDIKTLRVAKAIYKR 60
                                                                                                                                                                                                                                                                                                                                     Query Match 26.5%; Score 169.5; DB 2; Length 120; Best Local Similarity 33.1%; Pred. No. 2e-13; Matches 40; Conservative 23; Mismatches 55; Indels 3
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/890,865A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
APPLICANT: Constantini, Franklin
APPLICANT: Zeng, Li
TITLE OF INVENTION: AXIN GENE AND USES THEREOF
NUMBER OF SEQUENCES: 23
CORRESPONDENCE ADDRESS:
ATTORNEY/AGENT INFORMATION:
NAME: White, John
PRESISTRATION NUMBER: 28,678
REPERENCE/DOCKET NUMBER: 0575/54249
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 278-0400
TELEPERAX: (212) 391-0526
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADDRESSEE: Cooper & Dunham LLP
STREET: 1185 Avenue of the Americas
CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0575/54249
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CLASSIFICATION: 435
TORNEY A PROPERTY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 14
US-08-890-865A-11
; Sequence 11, Application US/08890865A
; Patent No. 6307019
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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TELECOMMUNICATION INFORMATION:
                                                                                                                        TELEPAX: (212)391-0526
INFORMATION FOR SEQ ID NO: 13:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             INFORMATION FOR SEQ ID NO: 11:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ATTORNEY/AGENT INFORMATION:
NAME: White, John P
REGISTRATION NUMBER: 28,6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (212) 278-0400
                                                                                                                                                                                         LENGTH: 120 amino acids
TYPE: amino acid
STRANDEDNESS: single
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                                                                                                                                                                                                                                                      TOPOLOGY: linear MOLECULE TYPE: Protein
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Sequence 918, Application US/09949016

Batent No. 6812339

GENERAL INFORMATION:
APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
FILE REFERENCE: CLOOL01307

CURRENT PELLING DATE: 2000-04-14

PRIOR FILING DATE: 2000-10-20
PRIOR FILING DATE: 2000-10-20

PRIOR FILING DATE: 2000-10-03

PRIOR FILING DATE: 2000-10-03

PRIOR FILING DATE: 2000-09-08

NUMBER OF SEQ ID NOS: 207012

SOFTWARE: FastSEQ for Windows Version 4.0

SEQ ID NO 9918
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        61 YIENNSVVSKQLKPATKTYIRDGIKKQQIGSVMFDQAQTEIQAVMEENAYQVFLTSDIYL 120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 WTKSLHSLLGDQDGAYLFRTFLEREKCVDTLDFWFACNGFRQMNLKDTKTLRVAKAIYKR 60
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 27.0%; Score 172.5; DB 2; Best Local Similarity 32.5%; Pred. No. 6e-13; Matches 40; Conservative 24; Mismatches 56;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STREET: COOPER & Dunham LLP STREET: 1185 Avenue of the Americas CITY: New York STATE: New York COUNTRY: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US/08/890,865A FILING DATE: 10-JUL-1997 CLASSIFICATION: 435
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 13, Application US/08890865A
Patent No. 6307019
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APPLICANT: Constantini, Franklin
APPLICANT: Zeng, Li
APPLICANT: Zeng, Li
APPLICANT: CONSTANTION: AXIN GENE AND
NUMBER OF SEQUENCES: 23
CORRESPONDENCE ADDRESS:
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                            121 EYV 123
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508 DLI 510
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509 DLI 511
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ORGANISM: Human
                                                                                                                                                  US-09-949-016-9918
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2 WSBAFDELLASKYGLAAFRAFLKSEFCEENIEFWLACEDFKKTK-SPQKLSSKARKIYTD 60
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                                                                                     1 WTKSLHSLLGDQDGAYLFRTFLEREKCVDTLDFWFACNGFRQMNLKDTKTLRVAKAIYKR 60
                                                                                                                                                                      61 YIENNSVVSKQLKPATKTYIRDGIKKQQIGSVMFDQAQTEIQAVMEENAYQVFLTSDIY 119
                                                                                                                                                                                                 61 YIENNSVVSKOLKPATKTYIRDGIKKQQIGSVMFDQAQTEIQAVMEENAYQVFLTSDIY 119
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APPLICANT: Hillman, Jennifer L.
APPLICANT: Goli, Surya K.
TITLE OF INVENTION: A NOVEL REGULATOR OF CELL SIGNALLING
TITLE OF INVENTION: A NOVEL REGULATOR OF CELL SIGNALLING
NUMBER OF SEQUENCES: 5
CORRESPONDENCE ADDRESS:
ADDRESSEE: INCTE PHARMACEUTICALS, INC.
STREET: 3174 Porter Drive
CITY: Palo Alto
STRATE: CA
COUNTRY: US
ZIP: 94304
ZIP: 94304
ZIP: 94304
ZIP: BONEL FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        26.2%; Score 167.5; DB 1; Length 211; 32.8%; Pred. No. 7.6e-13; tive 20; Mismatches 57; Indels 3
  Length 120;
Query Match 26.2%; Score 167.5; DB 2; Length Best Local Similarity 32.8%; Pred. No. 3.6e-13; Matches 39; Conservative 20; Mismatches 57; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: Billings, Lucy J.
REGISTRATION NUMBER: 36,749
REFERENCE/DOCKET NUMBER: PF-0157 US
TELECOMONICATION INFORMATION:
TELEPHONE: 415-85-0555
TELEFROX: 415-845-4166
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: FASTEN: DOS
SOFTWARE: FASTEQ Version 2.0
CURRENT APPLICATION NUMBER: US/08/740
FILING NATE.
                                                                                                                                                                                                                                                                                                                           Sequence 4, Application US/08748483
Patent No. 5955314
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEX:
TELEX:
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
LENGTH: 211 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
IMMEDIATE SOURCE:
LIBRARY: GenBank
CLONE: 292037
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity 32.8%
Matches 39; Conservative
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PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                   RESULT 15
US-08-748-483-4
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Search completed: April 20, 2006, 15:32:31 Job time : 11.277 secs us-09-587-574-2.rapbm

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April 20, 2006, 15:57:53 ; Search time 42.7309 Seconds (without alignments) 1202.714 Million cell updates/sec
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(cgn2 = f/ptcdata/1/pubpaa/USO8 PUBCOMB.pep:*

(cgn2 = f/ptcdata/1/pubpaa/USO9 PUBCOMB.pep:*

(cgn2 = f/ptcdata/1/pubpaa/USO3 PUBCOMB.pep:*

(cgn2 = f/ptcdata/1/pubpaa/USO3 PUBCOMB.pep:*

(cgn2 = f/ptcdata/1/pubpaa/USIOB PUBCOMB.pep:*

(cgn2 = f/ptcdata/1/pubpaa/USIOB PUBCOMB.pep:*
GenCore version 5.1.7
Copyright (c) 1993 - 2006 Biocceleration Ltd.
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Maximum DB seq length: 200000000
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Description	Sequence 1797, Ap	8, Ar	34,	36,			670,	670,	670,	270,	848,	2, Ag	20,	2, Ag	107,	107,	13, 2	11,	4, 7	4, Ap	165	2, Apj	14,	28	3970
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SUMMARIES	US-10-723-860-1797	98-831-8	-786	US-10-786-720-36	-374	US-10-182-936A-91		-10	US-10-477-173-670	US-10-092-900A-270	US-09-867-550-848	US-09-894-749-2	US-10-258-371B-20	US-10-989-054-2	US-10-087-684-107	US-10-218-779-107	US-10-899-422-13	US-10-899-422-11	US-09-894-749-4	US-10-989-054-4	US-10-094-749-1650	US-10-113-794A-2	US-10-428-487-14	-10-258-371B-2	US-10-108-260A-3970
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& Query Match	97.7	65.7		62.8						62.8		6	29.5	٩.	œ	28.9	28.4	28.4	7.	۲.	27.2	7.	٠	27.0	27.0
Score	624	420	401	401	401	401	401	401	401	401	186.5		186.5					•	177.5			172.5	172.5		172.5
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Sequence 1728, Apple Sequence 15, Apple Sequence 24, Appli Sequence 24, Appli Sequence 1507, Apple Sequence 1, Appli Sequence 1, Appli Sequence 339, Apple Sequence 176, Apple Sequence 1292, Apple Sequence 152, Apple Sequen	issue Sarcoma, Compositions &	Length 843;  Indels 0; Gaps 0;  FROMNLKDTKTLRVAKAIYKR 60	IAGNOSING AND TREATING COLON
4 US-10-087-192-1728 1 3 US-10-487-092-15-1728 1 4 US-10-258-3718-24 1 4 US-10-258-3718-24 1 4 US-10-408-765A-493 2 US-09-925-300-1507 3 US-09-925-300-1507 4 US-10-13-794A-1 5 3 US-09-736-457-339 5 3 US-09-962-941-339 5 3 US-09-962-941-339 6 4 US-10-017-754-339 7 US-10-13-872-339 8 4 US-10-283-017-339 9 US-10-283-017-339 1 US-09-925-301-1292 1 US-09-925-301-1292	alignments  ion US/10723860 1253606A1  tha  wendy M. Thords of Diagnosis of Soft Tilethods for Screening for Soft O193 NPUSO. 1203-11-26 1813 1010/10723,860 1 1203-11-26 1813 1310 1310 1813 1310 1310 1813 1310 1310 1813 1	Score 624; DB 5; Pred. No. 3.46-60; 4; Mismatches 1; RTFLEREKCVDTLDFWFACNG	tion US/10751736 40265230A1 , Robert ugene COMPOSITIONS AND METHODS FOR DIAGNOSING AND TREATING
28 172.5 27.0 776 29 172.5 27.0 917 31 167.5 26.2 211 32 167.5 26.2 211 33 167.5 26.2 211 34 167.5 26.2 211 35 167.5 26.2 211 36 165 25.8 196 37 165 25.8 196 41 165 25.8 196 42 165 25.8 196 44 165 25.8 196 45 165 25.8 196 46 165 25.8 196 47 165 25.8 196 48 165 25.8 196 49 165 25.8 196 49 165 25.8 196 49 165 25.8 196 41 165 25.8 196 42 165 25.8 196	RESULT 1  US-10-723-860-1797 ; Sequence 1797, Application US/10723860 ; Publication No. US20040253606A1 ; GENERAL INFORMATION: ; APPLICANT: A212, Natasha ; APPLICANT: Ginsburg, Wendy M. ; TTLE OF INVENTION: Methods of Diagnos: ; TTLE OF INVENTION: Methods for Screet; ; CURRENT FILING DATE: 2003-11-26 ; NUMBER OF SEQ ID NOS: 8393 ; SEQ ID NO 1797 ; LENGTH: 843 ; TTPE: PRT ; ORGANISM: Homo sapiens	Query Match         97.7%;           Best Local Similarity         95.9%;           Matches 118;         Conservative           Qy         1 WTKSLHSLIGDODGAYLF           Db         78 WTKSLHSLIGDODGAYLF           Qy         61 YIENNSVVSKQLKPATKT           Db         138 YIENNSIVSKQLKPATKT           Qy         121 EXV 123           Qy         121 EXV 123           Db         198 EXV 200	RESULT 2 US-10-751-736-116 Sequence 116, Application US/ Publication No. US20040265230 GENERAL INFORMATION: APPLICANT: Wacth APPLICANT: Martinez, Robert APPLICANT: Brown, Eugene APPLICANT: Liu, Wei TITLE OF INVENTION: COMPOSIT

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US-09-198-831-8
US-09-198-831-8
Sequence 8, Application US/09798831
Sequence 8. Application US/09798831
Sequence 8. Application US/09798831
Sequence 8. Application
GENERAL INFORMATION:
APPLICANT: KLEIN, Peter S.
TITLE OF INVENTION: SYNTHASE KINASE 3(BETA) ACTIVITY AND ACTIVATORS OF WNT TITLE OF INVENTION: SYNTHASE KINASE 3(BETA) ACTIVITY AND ACTIVATORS OF WNT TITLE OF INVENTION: SIGNALING
FILE REFERENCE: 2009-56.0391/306U1
CURRENT FILING DATE: 2000-03-01
PRIOR PILING DATE: 2000-03-01
NUMBER OF SEQ ID NOS: 12
                                                                                                                                                                                                                                                                                                                          61 YIENNSVVSKQLKPATKTYIRDGIKKQQIGSVMFDQAQTEIQAVMEENAYQVFLTSDIYL 120
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 WIKSLHSLLGDQDGAYLFRIFILEREKCVDTLDFWFACNGFROMNLKDTKT---LRVAKAI
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TITLE OF INVENTION: CANCERS
FILE REPERCEC: AM100927 (031896-002000)
CURRENT APPLICATION UNMBER: US/10/751,736
CURRENT FILING DATE: 2003-01-06
PRIOR APPLICATION NUMBER: US Provisional Application 60/438,000
PRIOR FILING DATE: 2003-01-06
NUMBER OF SEQ ID NOS: 54873
SOFTWARE: Patentin version 3.2
SEQ ID NO 116
LENGTH: 843
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; Score 420; DB 3; Length 842;
; Pred. No. 1.6e-37; ...
25; Mismatches 20; Indels
                                                                                                                                                                                                                                        Length 843;
                                                                                                                                                                                                                                                                        1; Indels
                                                                                                                                                                                                                                       Score 624; DB 5;
Pred. No. 3.4e-60;
                                                                                                                                                                                                                                                                       4; Mismatches
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61.1%;
                                                                                                                                                                                                                                     Query Match
Best Local Similarity 95.9%;
Matches 118; Conservative
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Best Local Similarity 61.18
Matches 77; Conservative
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US-09-798-831-8
                                                                                                                                                                                     ORGANISM: Homo sapiens
US-10-751-736-116
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SEQ ID NO 8
LENGTH: 842
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; Sequence 34, Application US/10786720

US-10-786-720-34

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Sequence 36, Application US/10786720;
Publication No. US20040191818A1
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Wyeth
APPLICANT: O'Toole, Margot
APPLICANT: O'Toole, Margot
APPLICANT: O'Toole, Margot
APPLICANT: Diu, Wei
APPLICANT: Liu, Wei
APPLICANT: Diu, Wei
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR DIAGNOSING AND TREATING AUTOIMMUNB
TITLE OF INVENTION: DISBASES
FILE REFERENCE: 031896-023000 (AM101331L)
CURRENT APPLICATION WUMBER: US/10/786,720
CURRENT FILING DATE: 2004-02-26
NUMBER OF SEQ ID NOS: 21135
SOFTWARE: Patentin version 3.2
                                                                                   APPLICANT: Liu, Wei
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR DIAGNOSING AND TREATING AUTOIMMUNE
TITLE OF INVENTION: DISEASES
TITLE OF INVENTION: DISEASES
CURRENT APPLICATION NUMBER: US/10/786,720
CURRENT APPLICATION NUMBER: 204-02-26
CURRENT FILING DATE: 2004-02-26
SOFTWARE: Patentin version 3.2
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                                                                                                                                                                                                                                                                                                                                                                                              62.0%; Score 401; DB 4; Length 461; 62.7%; Pred. No. 9.7e-36; Live 17; Mismatches 26; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; Score 401; DB 4; Length 826;
; Pred. No. 2e-35;
17; Mismatches 26; Indels
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62.7%;
Publication No. US20040191818A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 62.8%
Best Local Similarity 62.7%
Matches 79; Conservative
                                                                 O'Toole, Margot
                                                                                                                                                                                                                                                                                                                                                                                                                                               79; Conservative
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CORGANISM: Homo sapiens
US-10-786-720-36
                                                                                                                                                                                                                                                                                                            TYPE: PRT
ORGANISM: Homo sapiens
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Matches 79; Conserv
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                                         APPLICANT: Wyeth
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US-10-786-720-36
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                                                                                                                                                                                                                                                                     SEQ ID NO 34
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WS-10-182-936A-91

Sequence 91, Application US/10182936A

Publication No. US20040038860A1

Sequence 91, Application US/10182936A

Publicant USCOMATION:
APPLICANT: Allen, Kristina M.
APPLICANT: Bhat, Bheem
APPLICANT: Damagnez, Veronique
APPLICANT: Paworsky, Paul
TITLE OF INVENTION: Reagents and Method for Modulating DKK-Mediated Interactions
TITLE OF INVENTION: Reagents us (10/182,936A)
CURRENT APPLICATION NUMBER: US/10/182,936A

CURRENT FILING DATE: 2002-08-02
PRIOR APPLICATION NUMBER: US 60/291,311
PRIOR PILING DATE: 2001-05-17
PRIOR APPLICATION NUMBER: US 60/353,058

PRIOR FILING DATE: 2001-05-17

PRIOR FILING DATE: 2002-05-10

8 YKRYI-ENNSVVSKQLKPATKTYIRDGIKKQQIGSVMFDQAQTBIQAVMEENAYQVFLTS 116
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Sequence 670, Application US/10477238A

Publication No. US20040221326A1

GENERAL INFORMATION:
APPLICANT: Babij, Philip

APPLICANT: Waworsky, Paul

APPLICANT: Bodine, Peter Van Nest

TITLE OF INVENTION: Transgenic Animal Model of Bone Mass Modulation
FILE REFERENCE: 032796-212

CURRENT FILING DATE: 2003-11-10

PRIOR PAPLICATION NUMBER: US/10/477,238A

CURRENT FILING DATE: 2001-05-11

PRIOR PLLING DATE: 2001-05-11

PRIOR APPLICATION NUMBER: US 60/290,071
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 WIKSLHSLLGDQDGAYLFRTFLEREKCVDTLDFWFACNGFRQMNLKDT---KTLRVAKAI
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SOGTWARR: FastSEQ for Windows Version 4.0
SEQ ID NO 91
LENGTH: 900
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Best Local Similarity 62.7%
Matches 79; Conservative
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US-10-182-936A-91
                                                                                          DIYLEY 122
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                                                                                                         APPLICANT: Wyeth
APPLICANT: Wyeth
APPLICANT: Liu, Weith
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR DIAGNOSING AND TREATING AUTOIMMUNE
TITLE OF INVENTION: DISEASES
FILE REPERENCE: 031896-023000 (AMI01331L.)
CURRENT FILING DATE: 2004-02-26
NUMBER OF SEQ ID NOS: 21135
SOFTWARE: Patentin version 3.2
SOFTWARE: Patentin version 3.2
LENGTH: 862
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Publication No. US20030219793A1

GENERAL INFORMATION:

TILLE OF INVENTION: THE HIGH BONE MASS GENE OF 11q13.3

FILLE REFERENCE: 032796-021

CURRENT APPLICATION NUMBER: US/0/374,979

CURRENT FILING DATE: 2003-03-04

PRIOR FILING DATE: 2000-04-05

PRIOR FILING DATE: 2000-04-05

PRIOR FILING DATE: 1999-01-13

PRIOR FILING DATE: 1999-01-13

PRIOR FILING DATE: 1999-01-13

PRIOR FILING DATE: 1998-01-13

PRIOR FILING DATE: 1998-10-23

SEQ ID NO 91

LENGTH: 200
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b; Pred. No. 2.1e-35;
17; Mismatches 26
                 IS-10-786-720-35
Sequence 35, Application US/10786720
Publication No. US20040191818A1
GENERAL INFORMATION:
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Matches 79, Conservative
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Best Local Similarity 62.7%
Matches 79; Conservative
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ORGANISM: Homo sapiens
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US-10-374-979-91
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us-09-587-574-2.rapbm

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58 YKRYI-ENNSVVSKQLKPATKTYIRDGIKKQQIGSVMFDQAQTEIQAVMEENAYQVFLTS 116
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CURRENT FILING DATE: 2003-11-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 WIKSLHSLIGDODGAYLFRIFLEREKCVDILDFWFACNGFROMNLKDT---KTLRVAKAI
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                                                                                                                                                                                                                             APPLICANT: Genome Therapeutics Corporation and APPLICANT: Allen, Kristina M. APPLICANT: Yaworsky, Paul APPLICANT: Morales, Arturo J. APPLICANT: Anisowicz, Anthony APPLICANT: Liu, Wei
                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURKENT FILING DATE: 2003-11-10

PRIOR APPLICATION NUMBER: US 60/290,071

PRIOR FILING DATE: 2001-05-11

PRIOR FILING DATE: 2001-05-11

PRIOR FILING DATE: 2001-05-17

PRIOR FILING DATE: 2001-05-17

PRIOR FILING DATE: 2002-02-01

PRIOR FILING DATE: 2002-03-04

NUMBER OF SEQ ID NOS: 1086

SOFTWARE: FASELSEQ FOR WINDOWS VERSION 4.0

SEQ ID NO 670
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 270, Application US/10092900A publication No. US20040043382A1 GENERAL INFORMATION:
                                                                                                                                                               ; Sequence 670, Application US/10477173; Publication No. US20050070699A1; GENERAL INFORMATION:
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APPLICANT: Spytek, Kimberly A.
APPLICANT: Shenoy, Suresh G.
APPLICANT: Taupier Jr., Raymond J.
APPLICANT: Pena, Carol E.A.
APPLICANT: Li, Li
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Ramesh
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Gangolli, Esha A.
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Gusev, Vladimir
Ji, Weizhen
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APPLICANT: Babilion:
APPLICANT: Aworsky, Paul
APPLICANT: Babilion:
APPLICANT: Bodine, Peter Van US
APPLICANT: Bodine, Peter Van Nest
TILE BODINE: 03.796-179
CURRENT APPLICATION: Transgenic Animal Model of Bone Mass Modulation
FILE REFERENCE: 03.796-179
CURRENT APPLICATION NUMBER: US/10/680,287A
CURRENT FILING DATE: 2003-10-08
PRIOR PILING DATE: 2003-10-08
PRIOR PILING DATE: 2001-05-11
PRIOR PELICATION NUMBER: US 60/290,071
PRIOR PELICATION NUMBER: US 60/291,311
PRIOR PILING DATE: 2001-05-17
PRIOR PILING DATE: 2001-05-17
PRIOR PILING DATE: 2002-02-01
PRIOR PILING DATE: 2002-02-01
PRIOR FILING DATE: 2002-02-01
PRIOR FILING DATE: 2002-02-01
PRIOR FILING DATE: 2002-02-01
PRIOR APPLICATION NUMBER: US 60/353,058
PRIOR FILING DATE: 2002-02-01
PRIOR FILING DATE: 2002-02-01
PRIOR FILING DATE: 2002-03-04
NUMBER OF SEQ ID NOS: 812
SOFTWARE: PASLSEQ for Windows Version 4.0
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; PRIOR FILING DATE: 2001-05-17; PRIOR APPLICATION NUMBER: US 60/353,058; PRIOR FILING DATE: 2002-02-01; PRIOR PILING DATE: 2002-03-04; PRIOR FILING DATE: 2002-03-04; NUMBER OF SEQ ID NOS: 812; SOFTWARE: FastSEQ for Windows Version 4.0; SCQ ID NO 670; LENGTH: 900
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Application US/10680287A o. US20040244069A1
                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity 62.7*
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Best Local Similarity 62.78
Matches 79; Conservative
                                                                                                                                                                                                                                    ORGANISM: Homo sapiens
US-10-477-238A-670
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US-10-680-287A-670
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4; Gaps

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APPLICANT: Topper, James
TITLE OF INVENTION: No. US20020082206Alel Polynucleotides from Atherogenic Cells an
TITLE OF INVENTION: Thereby
TITLE OF INVENTION: Thereby
TITLE OF INVENTION: Thereby
FILE REFERENCE: 21402-013 (Cura-313)
CURRENT APPLICATION WUMBER: US/09/867,550
PRIOR APPLICATION NUMBER: USSN 60/208,427
PRIOR APPLICATION NUMBER: USSN 60/208,427
PRIOR APPLICATION NUMBER: USSN 60/208,427
STROID NOS: 2125
SOFTWARE: FastSEQ for Windows Version 4.0
STROID NO 848
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               55 KAIYKRYIENNSVVSKQLKPATKTYIRDGIKKQQIGSVMFDQAQTEIQAVMEBNAYQVFL 114
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Patent No. US20020081683A1

APPLICANT: Hodge, Martin R.

APPLICANT: Yowe, David

TITLE OF INVENTION: RESC. Containing Molecules and Uses Thereof
FILE REPERENCE: 5800-19, 035800/174680

CURRENT FILING DATE: 2001-06-27

PRIOR APPLICATION NUMBER: 09/244,314

PRIOR PILING DATE: 1999-02-04
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 186.5; DB 3
Pred. No. 2.9e-12;
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1 Similarity 34.6%;
44; Conservative 25
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SOFTWARE: Patentin Ver. 2.0
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ORGANISM: Homo sapiens
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Best Local Similarity
Matches 44; Conserva
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Best Local Similarity
Matches 44; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                           RPPLICANT: Burgess, Catherine E. TILLE OF INVENTION: No. US20040043382Alel Proteins and Nucleic Acids Encoding Same FILE REPERENCE: 21402-290C
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NUMBER OF SEQ ID NOS: 768
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Pred. No. 2.3e-35;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT AFFLICATION NUMBER: US/10/092, 900A
PRIOR APPLICATION NUMBER: USSN 60/274,322
PRIOR FILING DATE: 2001-03-08
PRIOR FILING DATE: 2001-03-08
PRIOR FILING DATE: 2001-04-13
PRIOR FILING DATE: 2001-04-13
PRIOR FILING DATE: 2001-03-08
PRIOR FILING DATE: 2001-03-08
PRIOR FILING DATE: 2001-03-08
PRIOR PLING DATE: 2001-03-08
PRIOR APPLICATION NUMBER: USSN 60/274,191
PRIOR APPLICATION NUMBER: USSN 60/279,995
PRIOR PLING DATE: 2001-03-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION NUMBER: US/10/092,900A
CURRENT FILING DATE: 2002-03-07
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Patent No. US20020082206Al
GENERAL INFORMATION:
APPLICANT: Leach, Martin D.
APPLICANT: Mehraban, Fuad,
              Vernet, Corine A.M.
Guo, Xiaojia Sasha
Tchernev, Velizar T.
Fernandes, Elma R.
                                                                                                    Casman, Stacie J.
Malyankar, Uriel M.
Gerlach, Valerie
                                                                                                                                                                                                   Anderson, David W.
                                                                                                                                                                                                                                                                                                                  Alsobrook, John P.
Lepley, Denise M.
                                                                                                                                                                                                                                               Catterton, Elina
Leite, Mario W.
                                                                                                                                                                                                                                                                                                                                                                   Daniel R
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Best Local Similarity 62.74
Matches 79; Conservative
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US-10-092-900A-270
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US-09-867-550-848
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us-09-587-574-2.rapbm

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RESULT 15
US-10-258-371B-20
| Sequence 20, Application US/10258371B |
| Sequence 20, Application US/10258371B |
| Publication No. US20040067903A1 |
| GENERAL INFORMATION: |
| APPLICANT: WILLIAMS-GAGNON, Alison |
| APPLICANT: WILLIAMS-GAGNON, Alison |
| TITLE OF INVENTION: NUCLEC ACIDS ENCODING A NOVEL REGULATOR OF G PROTEIN SIGNALING, |
| TITLE OF INVENTION: RGS18, AND USES THEREOF |
| TITLE OF INVENTION: NUCLEC ACIDS ENCODING A NOVEL REGULATOR OF G PROTEIN SIGNALING, |
| FILE REFERENCE: A3656 US PCT |
| CURRENT FILING DATE: 2000-06-02 |
| PRIOR FILING DATE: 2000-08-02 |
| PRIOR FILING DATE: 2000-08-02 |
| PRIOR FILING DATE: 2000-04-28 |
| NUMBER OF SEQ ID NOS: 38 |
| SOFTWARE: Patentin version 3.2 |
| LENGTH: 235 |
| LENGTH: 235
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ORGANISM: Homo sapiens
US-10-258-371B-20
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194 KSDIYLD 200
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Sequence 2, Appli
Sequence 4, Appli
Sequence 3946, Ap
Sequence 3945, Ap
Sequence 46, Appl
Sequence 7120, Ap
Sequence 20062, A
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48, Appl
27, Appl
3278, Ap
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25, Appl
3947, Ap
1069, Ap
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(without alignments)
794.148 Million cell updates/sec
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                                                                                                                                         1 WTKSLHSLLGDQDGAYLFRT......VMEENAYQVFLTSDIYLEYV 123
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1: /SIDSS/ptcdata/2/pubpaa/USO8 NEW PUB.pep:*

2: /SIDSS/ptcdata/2/pubpaa/USO6 NEW PUB.pep:*

3: /SIDSS/ptcdata/2/pubpaa/USO7 NEW PUB.pep:*

4: /SIDSS/ptcdata/2/pubpaa/PCT NEW PUB.pep:*

5: /SIDSS/ptcdata/2/pubpaa/USO9 NEW PUB.pep:*

6: /SIDSS/ptcdata/2/pubpaa/USO0 NEW PUB.pep:*

7: /SIDSS/ptcdata/2/pubpaa/US10 NEW PUB.pep:*

8: /SIDSS/ptcdata/2/pubpaa/US11 NEW PUB.pep:*
GenCore version 5.1.7
Copyright (c) 1993 - 2006 Biocceleration Ltd
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US-10-501-035-208
US-11-137-131-2
US-11-137-131-2
US-11-137-131-3
US-11-096-56BA-3946
US-11-096-56BA-3946
US-11-098-686-10183
US-11-098-686-10183
US-11-108-298-20062
US-11-108-298-20062
US-11-108-298-20062
US-11-108-298-20063
US-11-108-288-3947
US-10-10-65-56BA-3947
US-11-10-65-56BA-3947
US-11-10-65-56BA-3947
US-11-10-65-56BA-3947
US-11-10-65-56BA-3947
US-11-10-65-56BA-3947
US-11-10-65-56BA-3985
US-11-10-65-56BA-3089
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Maximum Match 100%
Listing first 45 summaries
                                             - protein search, using sw model
                                                                    April 20, 2006, 16:00:23
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Gapop 10.0 , Gapext 0.5
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Match
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Maximum DB
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Sequence 215, Application US/10501035
Publication No. US20060046249A1
GREEAL INFORMATION:
GREEAL INFORMATION:
GREEAL INFORMATION:
TITLE OF INVENTION: ACTIVITY OF COMPOUNDS THAT INTERACT WITH PROTEIN TYROSINE KINATITLE OF INVENTION: AND/OR PROTEIN TYROSINE KINASE PATHWAYS
FILE REFERENCE: DO185 PCT
CURRENT APPLICATION NUMBER: US/10/501,035
FILE REFERENCE: 2004-07-09
PRIOR APPLICATION NUMBER: US 60/350,061
PRIOR PILING DATE: 2002-01-18
NUMBER OF SEQ ID NOS: 795
SEQ ID NO 215
LENGTH: 900
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27553, A
27552, A
31249, A
31248, A
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12890, A
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28742,
28741,
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US-11-188-298-2708

US-10-508-263-1056

US-10-508-263-1066

US-10-467-657-1604

US-10-143-984A-40

US-10-10-79-463-8912

US-11-096-568A-27554

US-11-096-568A-27553

US-11-096-568A-21249

US-11-096-568A-31249

US-11-096-568A-28741
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Matches 79; Conservative
    995
168
156
663
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718
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378
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US-10-501-035-208

US-10-501-035-208

Sequence 208, Application US/10501035

Publication No. US20060046249A1

GENERAL INFORMATION:

APPLICANT: Bristol-Myers Squibb Company

TITLE OF INVENTION: IDENTIFICATION OF POLYNUCLEOTIDES AND POLYPEPTIDE FOR PREDICTING

TITLE OF INVENTION: ACTIVITY OF COMPONDES THAT INTERACT WITH PROFEIN TYROSINE KINA

TITLE OF INVENTION: AND/OR PROFEIN TYROSINE KINASE PATHWAYS

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10988, A

Sequence Sequence

Sequence

66.5 66.5 66

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56 AIYKRYIENNSVVSKQLKPATKTYIRDGİKKQQIĞSVM---FDQAQTEIQAVMEENAYQV 112
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            377 TVYLADILFCESALFYFSEYMEKEDAVNILQFWLAADNFQSQLAAKKGQYDGQEAQNDAM 436
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 74.5; DB 7;
Pred. No. 5.7;
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APPLICANT: Braun, A.

TITLE OF INVENTION: POLYMORPHIC KINASE ANCHOR PROTICE OF INVENTION: POLYMORPHIC ACIDS ENCODING THE STILE REFERENCE: 24736-2035

CURRENT APPLICATION NUMBER: US/11/137,131

CURRENT APPLICATION NUMBER: US/09/834,700

PRIOR APPLICATION NUMBER: 60/217,251

PRIOR FILING DATE: 2000-07-10

PRIOR FILING DATE: 2000-07-10

PRIOR FILING DATE: 2000-07-10

PRIOR PLING DATE: 2000-10-13

NUMBER OF SEQ ID NOS: 25

SOFTWARE: FASTSEQ for Windows Version 4.0
PRIOR APPLACE.
PRIOR FILING DATE: ZUUE
PRIOR FILING DATE: ZUUE
PRIOR FILING DATE: 2000-07-10
PRIOR FILING DATE: 2000-10-13
PRIOR FILING DATE: 2000-10-13
NUMBER OF SEQ ID NOS: 25
NUMBER OF SEQ ID NOS: 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   25; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity 21.4%;
Matches 28; Conservative 2
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Best Local Similarity 21.4%
Matches 28; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                        ORGANISM: Homo Sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                          US-11-137-131-2
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TITLE OF INVENTION: IDENTIFICATION OF POLYNUCLEOTIDES FOR PREDICTING ACTIVITY OF
TITLE OF INVENTION: COMPOUNDS THAT INTERACT WITH AND/OR MODULATE PROTEIN TYROSINE
TITLE OF INVENTION: KINASES AND/OR PROTEIN TYROSINE KINASE PATHWAYS IN LUNG CANCER
TITLE OF INVENTION: CELLS
FILE REPERENCE: 10001 NP
CURRENT APPLICATION NUMBER: US/11/169,041
CURRENT FILING DATE: 2005-06-28
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              61 YIENNSVVSKOLKPATKTYIRDGIKKQOIGSVMFDQAQTEIQAVMEENAYQVFLTSDIY 119
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Publication No. US20060003358A1
GENERAL INFORMATION:
APPLICANT: Braun, A.
TITLE OF INVENTION: POLYMORPHIC KINASE ANCHOR PROTEINS AND
TITLE OF INVENTION: NUCLEIC ACIDS ENCODING THE SAME
FILE REFERENCE: 24736-2035
CURRENT APPLICATION NUMBER: US/11/137,131
CURRENT FILING DATE: 2005-05-24
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Local Similarity 32.8%; Pred. No. 1.1e-10;
Les 39; Conservative 20; Mismatches 57
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20; Mismatches
                   ; FILE REPERENCE: D0185 PCT
; CURRENT APPLICATION NUMBER: US/10/501,035
; CURRENT FILING DATE: 2004-07-09
; PRIOR APPLICATION NUMBER: US 60/350,061
; PRIOR FILING DATE: 2002-01-18
; NUMBER OF SEQ ID NOS: 795
; SOFTWARE: PatentIn version 3.2
; SOFTWARE: Patent 218
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PRIOR APPLICATION NUMBER: 60/584,405
PRIOR FILING DATE: 2004-06-30
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SOFTWARE: PatentIn version 3.2
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US-11-169-041-234
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Matches 39, Conserv
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LENGTH: 211
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Matches
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11; Gaps

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64 GYLLFREFCLNHMERAKPLVEFYDBIKKYBKLDSEBERTVK-SRBIFDLYIMKELLSCSH 122
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Kapur, Vivek and Gebhart, Connie J.
TITLE OF INVENTION: FROM LAWSONIA INTRACELLULARIS AND METHODS OF USING
TITLE OF INVENTION: FROM LAWSONIA INTRACELLULARIS AND METHODS OF USING
FILE REFERENCE: 09531-128001
CURRENT APPLICATION NUMBER: US/11/098,686
CURRENT FILING DATE: 2005-04-04
PRIOR FILING DATE: 2003-10-01
PRIOR FILING DATE: 2003-10-01
PRIOR FILING DATE: 2003-10-04
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                TITLE OF INVENTION: Polypeptides and Nucleic Acids Encoding Same TITLE REPERENCE: 21402-225
CURRENT APPLICATION: Polypeptides and Nucleic Acids Encoding Same FILE REFERENCE: 21402-225
CURRENT FILING DATE: 2005-04-21
PRIOR APPLICATION NUMBER: 60/256,704
PRIOR PLING DATE: 2000-12-19
PRIOR FILING DATE: 2000-12-19
PRIOR FILING DATE: 2000-12-19
PRIOR FILING DATE: 2000-12-20
PRIOR PLING DATE: 2000-12-20
PRIOR PRIOR DATE: 2001-08-10
PRIOR PLING DATE: 2001-08-10
PRIOR PLING DATE: 2001-08-10
PRIOR PLING DATE: 2001-08-10
PRIOR PLING DATE: 2001-09-19
PRIOR PLING DATE: 2001-09-29
PRIOR PLING DATE: 2001-09-14
PRIOR PLING DATE: 2001-09-14
PRIOR PLING DATE: 2001-09-14
PRIOR PLING DATE: 2001-09-14
PRIOR PLING DATE: 2001-09-14
PRIOR PLING DATE: 2001-05-14
PRIOR PLING DATE: 2001-05-14
PRIOR PLING DATE: 2001-05-02
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Best Local Similarity 20.4%; Pred. No. 14;
Matches 23; Conservative 33; Mismatches 5:
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22.2%; Pred. No. 16;
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SOFTWARE: PastSEQ for Windows Version 4.0
SEQ ID NO 10183
LENGTH: 691
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Publication No. US20060024696A1
GENERAL INFORMATION:
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SEQ ID NO 46
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Best Local Similarity
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                                                                                                  APPLICANT: Alexandrov, Nickolai et al.
TITLE OF INVENTION: Sequence-Determined DNA Fragments and Corresponding Polypeptides
TITLE OF INVENTION: Therby
FILLE OF INVENTION: Therby
FILLE OF INVENTION: Therby
CURRENT APPLICATION NUMBER: US/11/096,568A
CURRENT APPLICATION NUMBER: US/11/096,568A
NUMBER OF SEQ ID NOS: 34471
SEQ ID NOS: 34471
LENGTH: 285
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TITLE OF INVENTION: Sequence-Determined DNA Fragments and Corresponding Polypeptides
TITLE OF INVENTION: Therby
FILE REFERENCE: 2750-15920032
CURRENT APPLICATION NUMBER: US/11/096,568A
CURRENT APPLICATION NUMBER: US/11/096,568A
NUMBER OF SEQ ID NOS: 34471
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          71 OLKPATKTYIRDGIKKQQIGSVMFDQAQTEIQAVMEENAYQVFLTS-DIY 119
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            69 DWNPSA-TSITVGLSDGSVSIVSFLESKLEIQEBWKAHDYELWTISFDIH 117
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        71 QLKPATKTYIRDGIKKQQIGSVMFDQAQTEIQAVMEENAYQVFLTS-DIY 119
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                                                                                                                                                                                                                                                                                                                                                                                                     | FEATURE:
| NAME/KEY: misc feature
| LOCATION: (1).7(285)
| JOHTER INFORMATION: Ceres Seq. ID no. 13594271
| US-11-096-568A-3946
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME/KEY: misc_feature
i LOCATION: (1)..(344)
cother information: Ceres Seq. ID no. 13594270
US-11-096-5688-3945
                      Sequence 3946, Application US/11096568A Publication No. US20060048240A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 3945, Application US/11096568A Publication No. US20060048240A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                               ORGANISM: Glycine max
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JS-11-096-568A-3946
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US-11-113-424-46
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Sequence 46, Application US/11113424

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                                               KRYIENNSVVSKOLKPATKTYIRDGIKKQQIGSVMFDQAQTEIQAVMEENAYQVFLTSDI 118
182 WAKN----IGFLDLAHGFSIAMKEE-----IDFKIEARNFEOVSDSLKNSETKVKIPKVY 232
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 14 GAYLFRIFL--EREKCVDILDFWFACNGFROMNLKDIKTLRVAKAIYKRYIENNSV-VSK 70
                                                                             233 KKYSNSKILVLEFL------DGVSVKS-GSALLNELQIDTKKVQR----QLF---DC
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                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Gangolli et al. TITLE OF INVENTION: Polypeptides and Nucleic Acids Encoding Same FILE REFERENCE: 21402-225
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 688;
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Sequence 25, Application No. US20060029983A1
GENERAL INFORMATION:
APPLICANT: OAKLEY, ROBERT H.
APPLICANT: HUSSON, CHRISTINE C.
TITLE OF INVENTION: CONSTITUTIVELY TRANSLOCATING CELL LINE
FILE REFERENCE: NRK.108
CURRENT FILLING NUMBER: US/11/040,218
CURRENT FILLING DATE: 2005-01-21
PRIOR APPLICATION NUMBER: US/10/788,197
PRIOR FILING DATE: 2004-02-26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         58; Indels
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18;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT PELLICATION NUMBER: US/11/113,424

CURRENT PELLING DATE: 2005-04-21

PRIOR PELLING DATE: 2005-04-21

PRIOR FILING DATE: 2000-12-19

PRIOR PELLING DATE: 2001-08-10

PRIOR APPLICATION NUMBER: 60/311,590

PRIOR APPLICATION NUMBER: 60/311,590

PRIOR PELLING DATE: 2001-08-10

PRIOR PELLING DATE: 2001-08-10

PRIOR FILING DATE: 2001-08-29

PRIOR FILING DATE: 2001-08-29

PRIOR PELLORTION NUMBER: 60/315,617

PRIOR PELLOR DATE: 2001-08-29

PRIOR APPLICATION NUMBER: 60/315,617

PRIOR APPLICATION NUMBER: 60/307,506

PRIOR APPLICATION NUMBER: 60/307,506

PRIOR PELLING DATE: 2001-07-24

PRIOR FILING DATE: 2001-07-24

PRIOR FILING DATE: 2001-07-24

PRIOR FILING DATE: 2001-08-14

PRIOR PELLORTION NUMBER: 60/294,075

PRIOR APPLICATION NUMBER: 60/294,075

PRIOR PELLOR DATE: 2001-06-29

PRIOR PELLOR DATE: 2001-06-29

PRIOR PELLOR DATE: 2001-06-29

PRIOR PELLOR DATE: 2001-06-29

PRIOR PELLOR DATE: 2001-06-29

PRIOR PELLOR DATE: 2001-06-29
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Pred. No. 1
                                                                                                                                                                                                                                                                                                                             Sequence 49, Application US/11113424 Publication No. US20050260713A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity 20.5%;
Matches 23; Conservative 2'
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                                                                                                                                                  119' YLEYV 123
                                                                                                                                                                                                  276 ILEQI 280
                                                                                                                                                                                                                                                                               RESULT 12
US-11-113-424-49
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LENGTH: 68
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TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO BACTEROIDES FRA

TITLE OF INVENTION: FOR DIAGNOSTICS AND THERAPEUTICS

FILE REFERENCE: PATHOG-03DIV2

CURRENT APPLICATION NUMBER: US/11/079,463

CURRENT PILING DATE: 2005-03-14

PRIOR APPLICATION NUMBER: US 60/128,705

PRIOR FILING DATE: 1999-04-09

PRIOR FILING DATE: 2000-04-04

NUMBER OF SEQ ID NOS: 10444
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; OTHER INFORMATION: Identity of amino acid sequences at the above locations are unknous-11-079-463-7120
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       KCVDTLDFWFACNGFRQMNLK--DTKTLRVAKAIYKRYIENNSVVSKQLKPATKTYIRDG 83
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; Publication No. US2066075522A1
; GENERAL INFORMATION:
    APPLICANT: Abad, Mark S. et al.
; TITLE OF INVENTION: GENES AND USES FOR PLANT IMPROVEMENT;
FILE REFERENCE: 38-21(53452)8
; CURRENT APPLICATION NUMBER: US/11/188,298
; CURRENT FILING DATE: 2005-07-22
; PRIOR APPLICATION NUMBER: 60/592,978
; PRIOR FILING DATE: 2004-07-31
; NUMBER OF SEQ ID NOS: 22569
; SEQ ID NO 20062
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    7; Length 430;
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14;
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                               541 YKKNDIYSETGISFEYTINDVLTIHARNCFTLHNMKNIF 679
                                                                                                       84 IKKOOIGS---VMFDQAQTEIQAVMEENAYQVFLTSDIY 119
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11.0%; Score 70; DB 'Best Local Similarity 24.0%; Pred. No. 14; Matches 30; Conservative 25; Mismatches
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US-11-188-298-20062
                                                                                                                                                                                                                                                                             Sequence 7120, Application US/11079463
Publication No. US20060073161A1
GENERAL INFORMATION:
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Best Local Similarity 25.0
Matches 30; Conservative
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; TITLE OF INVENTION: The Complete Genome and Protein Sequences of the Hyperthermophi; TITLE OF INVENTION: Methanopyrus Kandleri AU19 and Monophyly of Archael Methanogen; TITLE OF INVENTION: and Methods of Use Thereof TITLE OF INVENTION: and Methods of Use Thereof CURENT PERBRENCE: FIDOUL CURENT APPLICATION NUMBER: US/10/506,454

CURENT APPLICATION NUMBER: US/10/506,454

CURRENT FILING DATE: 2003-03-04

PRIOR FILING DATE: 2003-03-04

NUMBER OF SEQ ID NOS: 1722

SEQ ID NO 1069

LENGTHARE: Patentin version 3.2

LENGTH: 312

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; ORGANISM: Methanopyrus kandleri
US-10-506-454-1069
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Best Local Similarity 28.64
Matches 20; Conservative
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Publication No. US20060048240A1
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Alexandrov, Nickolai et al.
TITLE OF INVENTION: Sequence-Determined DNA Fragments and Corresponding Polypeptides
TITLE OF INVENTION: Therby
FILE REFERENCE: 2750-1592PUS2
CURRENT APPLICATION NUMBER: US/11/096,568A
CURRENT APPLICATION DATE: 2005-04-01
NUMBER OF SEQ ID NOS: 34471
SEQ ID NO 3947
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         64 GFLLFKDFCLNEINEAVPQVKFYEEIKEYEKLDNEEDRLCR-SRQIYDAYIMKELLSCSH 122
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              4; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         16; Mismatches 34; Indels 11; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           71 QLKPATKTYIRDGIKKQQIGSVMFDQAQTEIQAVMEENAYQVFLTSDIYLEY 122
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (23 PFSKQAVEHVQSHLSKKQVTSTLFQPYIEEICESLRGDIFQKFMESDKFTRF 174
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 247;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 688;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              58; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          10.9%; Score 69.5; DB 7; 27.4%; Pred. No. 5.6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
11.0%; Score 70; DB 7;
Best Local Similarity 20.5%; Pred. No. 18;
Matches 23; Conservative 27; Mismatches 56
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; LOCATION: (1)...(247)
; OTHER INFORMATION: Ceres Seq. ID no. 13594272
US-11-096-568A-3947
                                                                                                              PRIOR APPLICATION NUMBER: PCT/USO3/145E

PRIOR PELING DATE: 2003-05-12

PRIOR PLING DATE: 2003-05-13

PRIOR FILING DATE: 2002-05-13

PRIOR FILING DATE: 2002-06-07

PRIOR FILING DATE: 2002-08-07

NUMBER OF SEQ ID NOS: 94

SOGTWARE: PRICENTIN Ver. 3.2

SEQ ID NO 25

LENGTH: 688
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      97 AQTEIQAVMEENAYQVFLTS-DIY 119
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SKLEIQEEWKAHDYELWTTSFDIH 79
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; Sequence 1069, Application US/10506454
; Publication No. US20060068386Al
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Slesarev, Alexi I
APPLICANT: Mezhevaya, Katja V
APPLICANT: Polushin, Nikolai N
APPLICANT: Shcherbinina, Olga V
APPLICANT: Shakhova, Vera V
APPLICANT: Malykh, Andrei G
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 10.99
Best Local Similarity 27.45
Matches 23; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: PRT; ORGANISM: Homo sapiens
US-11-040-218-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME/KEY: misc feature
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ORGANISM: Glycine max
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Sequence 2, Appli Sequence 11780, A Sequence 12181, A Sequence 4711, Ap Sequence 2, Appli Sequence 2, Appli Sequence 2, Appli Sequence 2, Appli Sequence 2, Appli Sequence 613, Ap Sequence 613, Ap Sequence 613, Ap Sequence 613, Ap Sequence 613, Ap Sequence 613, Ap

Scoring table:

Searched:

Database :

Perfect score:

Sequence:

OM protein

Run on:

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Length 855;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY: New York

COUNTRY: US

COUNTRY: US

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

COMPUTER: IBM PC compatible

COMPUTER: IBM PC compatible

COMPUTER: PatentIn Release #1.0, Version #1.30

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/890,865A

FILING DATE: 10-JUL-1997

CLASSIFICATION: 435

ATTORNEY/AGENT INFORMATION:

NAME: White, John P

REGISTRATION NUMBER: 28,678

REGISTRATION NUMBER: 28,678

REGISTRATION NUMBER: 0575/54249

TELEFHONE: (212)278-0400

TELEFRONE: (212)278-0400

TELEFRAM: (212)278-0566

INFORMATION FOR SEQ ID NO: 10: SEQUENCE CHARACTERISTICS:

LEMGTH: 855 maino acids

LEMGTH: 855 maino acids

TVENEY.
                                                                                                                                                                                                                                                                                                                                        RESULT 1
US-08-890-865A-10
i Sequence 10, Application US/08890865A
j Patent No. 6307019
i GENERAL INFORMATION:
i APPLICANT: Constantini, Franklin
i APPLICANT: Zeng, Li
i TITLE OF INVENTION: AZIN GENE AND USES THEREOF
NUMBER OF SEQUENCES: 23
CORRESPONDENCE ADDRESS:
i CORRESPONDENCE ADDRESS:
i STREET: 1185 Avenue of the Americas
CITY: New York
STATE: New York
US-09-902-540-11516

US-08-936-135-4

US-08-936-135-4

US-09-248-796A-21780

US-09-902-540-16283

US-09-902-540-16283

US-09-107-522A-4711

US-08-981-215-1

US-08-981-215-1

US-09-910-215-1

US-09-940-921B-4

US-09-940-921B-2

US-09-940-921B-2

US-09-940-921B-2

US-09-940-921B-2

US-09-940-921B-2

US-09-940-921B-2

US-09-940-921B-2

US-09-940-921B-2

US-09-543-661A-4336

US-09-543-661A-4336

US-09-543-661A-4336
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Ouery Match 57.9%; Score 157.5; DB 2; Best Local Similarity 59.3%; Pred. No. 3.2e-11; Matches 32; Conservative 11; Mismatches 10;
   2584
25884
16584
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4739
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MOLECULE TYPE: Protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  single
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STRANDEDNESS:
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26 051, A
26 051, A
2, Appli
2, Appli
15206, A
13432, A
21807, A
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265, App
30322, A
2791, Ap
2, Appli
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                                                                                                                     (without alignments)
989.497 Million cell.updates/sec
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                                                                                                       April 20, 2006, 15:30:33 ; Search time 4.51187 Seconds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence Sequence
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1 ANGQVSLPHFPRTHRLPKEM......RLEKLKLELESRHSLEERLQ
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              GenCore version 5.1.7
Copyright (c) 1993 - 2006 Biocceleration Ltd
                                                                                                                                                                                                                                                                                                                                                                                                                                                     /cgn2_6/ptodata/1/iaa/5_COMB.pep:*
/cgn2_6/ptodata/1/iaa/6_COMB.pep:*
/cgn2_6/ptodata/1/iaa/H_COMB.pep:*
/cgn2_6/ptodata/1/iaa/FCTUS_COMB.pep:*
/cgn2_6/ptodata/1/iaa/RE_COMB.pep:*
/cgn2_6/ptodata/1/iaa/RE_COMB.pep:*
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US-08-890-865A-4
US-08-252-991A-26051
US-08-458-023B-2
US-09-111-556A-2
US-08-360-758-2
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-09-248-796A-21807
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US-09-577-304A-265
US-09-252-991A-30322
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-09-577-304A-268
-09-758-282B-261
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                                                                                                                                                                                                                                                                        572060 seqs, 82675679 residues
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                                                                                                                                                                                                                                                                                                                                                                            Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                                         - protein search, using sw model
                                                                                                                                                                                                                                          Gapop 10.0 , Gapext 0.5
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                                                                                                                                                                                                                                                                                                                                   Minimum DB seq length: 0
Maximum DB seq length: 2000000000
                                                                                                                                                                 US-09-587-574-3
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157.5 142.5 141.5 69.5

Result Š Gaps

1;

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Sequence 26051, Application US/09252991A

Patent No. 6551795

GENERAL INFORMATION:
APPLICANT: Marc J. Rubenfield et al.
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS TITLE OF INVENTION: AERGIGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 107196 136
CURRENT APPLICATION NUMBER: US/09/252,991A
CURRENT PILING DATE: 1999-02-18
PRIOR APPLICATION NUMBER: US 60/074,788
PRIOR APPLICATION NUMBER: US 60/074,788
PRIOR APPLICATION NUMBER: US 60/094,190
PRIOR APPLICATION NUMBER: US 60/094,190
PRIOR APPLICATION NUMBER: US 60/094,190
PRIOR APPLICATION NUMBER: US 60/094,190
PRIOR APPLICATION NUMBER: US 60/094,190
PRIOR APPLICATION NUMBER: US 60/094,190
PRIOR APPLICATION NUMBER: US 60/094,190
PRIOR APPLICATION NUMBER: US 60/094,190
PRIOR APPLICATION NUMBER: US 60/094,190
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    13; Indels 15;
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                                                                                                                                                                                                                                                                                                                                                                                                                      Length 900;
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; Pred. No. 0.78;
10; Mismatches 13; Indels 1:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
APPLICANT: Berka, Randy M.
APPLICANT: Yoder, Wendy
APPLICANT: Taggi, Shinobu
APPLICANT: Boominachan, Karuppan C.
TITLE OF INVENTION: ASPERGILLUS EXPRESSION SYSTEM
                                                                                                                                                                                                                                                                                                                                                                                                                      52.0%; Score 141.5; DB 2; 56.6%; Pred. No. 3.3e-09;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     8; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 2, Application US/08458023B Patent No. 5667990
                                                                  NAME: White, John P
REGISTRATION NUMBER: 28,678
REFERENCE/DOCKET NUMBER: 0575
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212)291-0526
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTER.ESTICS:
LENGRYH: 900 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ORGANISM: Pseudomonas aeruginosa
US-09-252-991A-26051
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          25.6%;
32.1%;
FILING DATE: 10-JUL-1997
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity 32.1<sup>3</sup>
Matches 18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity 56.6
Matches 30; Conservative
                                                                                                                                                                                                                                                                                                        single
                                                                                                                                                                                                                                                                                                                              TOPOLOGY: linear MOLECULE TYPE: Protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
ADDRESSEE: No. 56679
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 198 ANGELTLSH---
                                                                                                                                                                                                                                                                                                        STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-09-252-991A-26051
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQ ID NO 26051
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US-08-458-023B-2
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                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDLUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/890,865A
FILING DATE: 10-JUL-1997
CLASSIFICATION: 435
ATTORNEY/ABENT INFORMATION:
NAME: White, John P
REGISTRATION NUMBER: 28,678
REGISTRATION NUMBER: 28,678
REGISTRATION NUMBER: 0575/54249
TELECCOMMUNICATION INFORMATION:
TELEPHONE: (212)278-0400
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Patent No. 6307019
GENERAL INFORMATION:
FAPPLICANT: Constantini, Franklin
APPLICANT: Zeng, Li
TITLE OF INVENTION: AXIN GENE AND USES THEREOF
VUMBER OF SEQUENCES: 23
CORRESPONDENCE ADDRESS:
ADDRESSEE: Cooper & Dunham Lip
STREET: 1185 Avenue of the Americas
CITY: New York
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                                                                                                                                                                      AXIN GENE AND USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  8e-09;
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                                                                                                                                                                                STREET: 1185 Avenue of the Americas CITY: New York COUNTRY: US
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                             GENERAL INFORMATION:
APPLICANT: Constantini, Franklin
APPLICANT: Zeng, Li
TITLE OF INVENTION: AXIN GENE ANI
NUMBER OF SEQUENCES: 23
CORRESPONDENCE ADDRESS:
                                                Sequence 1, Application US/08890865A Patent No. 6307019
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INFORMATION FOR SEQ ID NO: 1:
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TYPE: amino acid
STRANDEDNESS: single
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Matches 30; Conservative
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SSEE: No. 60748630 No. 6074863disk of No. 6074863th America, Inc. 195 Lexington Avenue, 64th Floor New York
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Patent No. 6074863
GENERAL INFORMATION:
APPLICANT: Svendsen, Allan
APPLICANT: Borch, Michi
APPLICANT: Borch, Michi
APPLICANT: Clausen, Ib G
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APPLICANT: Clausen, 
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Pred. No. 5.3;
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COUNTRY: United States of America
ZID: 10174-6401
COMPUTER: Tape
COMPUTER: Tape
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/360,758
FILING DATE: 22-DEC-1994
CLASSIFICATION: 435.
PRIOR APPLICATION 135.
PRIOR APPLICATION NUMBER: DK PCT/DK93/00225
FILING DATE: 03-JUN-1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          336 ASYTVSVPKFPRFIWHAIPDEIVPYQPAA 364
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                                                                                                                                                                                                                                                                                                                                                                                                                         23.9%; Score 65; 48.3%; Pred. No. 9
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 212-867-0123
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                                                                                                                                  463 amino acids
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REGISTRATION NUMBER: 33
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Matches 14; Conservative
TELEFAX: 212-878-9655
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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INFORMATION FOR SEQ ID NO:
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                                                                                                                                                                                                                 single
                                                                                                                                                                                                                                                                                      MOLECULE TYPE: protein
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                                                                                                                                                                 TYPE: amino acid
STRANDEDNESS: sir
TOPOLOGY: linear
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Best Local Similarity
Matches 14; Conserva
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US-08-360-758-2
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APPLICANT: Clausen, Ib G
APPLICANT: Clausen, Mogens T
TITLE OF INVENTION: C. ANTARCTICA LIPASE AND LIPASE VARIANTS
WUMBER OF SEQUENCES: 11
CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                      SOFTWARE STATES TO SOFTWARE STATES SOFTWARE STATES SOFTWARE STATES SOFTWARE STATES SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE SOFTWARE S
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/111,556A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 65; DB 3
Pred. No. 5.3;
5; Mismatches
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APPLICATION NUMBER: DK PCT/DK93/00225
FILING DATE: 03-JUN-1993
ATTORNEY/AGENT INFORMATION:
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REFERENCE/DOCKET NUMBER: 3748.214-US
TELECOMMUNICATION INFORMATION:
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22-DEC-1994
                          COUNTY: 0.5A

ZIP: 10174-6201

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS
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Pathar, Shamkant A
Egel-Mitani, Michi
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INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
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Best Local Similarity 48.3%;
Matches 14; Conservative
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amino acid
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        USA
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US-09-248-796A-21807
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US-10-200-012-4
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                                                                                                                                                GENERAL INFORMATION:
APPLICANT: Goldman, Barry S.
APPLICANT: Hinkle, Gregory J.
APPLICANT: Hinkle, Steven C.
APPLICANT: Steven C.
APPLICANT: Steven C.
TITLE OF INVENTION: Myxococcus xanthus Genome Sequences and Uses Thereof FILE REFERENCE: 38-10(15849)B
CURRENT APPLICATION NUMBER: US/09/902,540
CURRENT FILING DATE: 2001-07-10
PRIOR FILING DATE: 2000-07-10
NUMBER OF SEQ ID NOS: 16825
SEQ ID NO 15206
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:

APPLICANT: Goldman, Barry S.

APPLICANT: Hinkle, Gregory J.

APPLICANT: Slater, Steven C.

APPLICANT: Wiegand, Roger C.

TITLE OF INVENTION: Myxococcus xanthus Genome Sequences and Uses Thereof FILE REFERENCE: 38-10(15499) B CURRENT APPLICATION NUMBER: US/09/902,540

CURRENT FILING DATE: 2001-07-10
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Pred. No. 7.8;
4; Mismatches
336 ASYTVSVPKFPRFIWHAIPDEIVPYQPAA 364
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 61;
Pred. No. 1
                                                                                                                  Sequence 15206, Application US/09902540 Patent No. 6833447
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION NUMBER: 60/217,883
PRIOR FILING DATE: 2000-07-10
NUMBER OF SEQ ID NOS: 16825
SEQ ID NO 14932
LENGTH: 188
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Best Local Similarity 39.5%;
Matches 17; Conservative
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Best Local Similarity 50.0%;
Matches 15; Conservative 4
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                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: PRT ORGANISM: Myxococcus xanthus
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GENERAL INFORMATION:
APPLICANT: Keith Weinstock et al
APPLICANT: Keith Weinstock et al
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO CANDIDA ALBIC;
TITLE OF INVENTION: FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 107196.132
CURRENT APPLICATION NUMBER: US 60/074,725
PRIOR FILING DATE: 1998-02-13
PRIOR FILING DATE: 1998-02-13
PRIOR FILING DATE: 1998-02-13
NUMBER OF SEQ ID NOS: 28208
SEQ ID NOS: 28208
SEQ ID NOS: 28208
Page

**APPLICANT: Slater, Steven C.

**APPLICANT: Wiegand, Roger C.

**TITLE OF INVENTION: Myxococcus xanthus Genome Sequences and Uses Thereof

**FILE REFERENCE: 38-10(15849)B

**CURRENT APPLICATION NUMBER: US/09/902,540

**CURRENT APPLICATION NUMBER: US/09/902,540

**CURRENT APPLICATION NUMBER: US/09/902,540

**PRIOR FILING DATE: 2001-07-10

**PRIOR FILING DATE: 2000-07-10

**NUMBER OF SEQ ID NOS: 16825

**SEQ ID NO 13429

**LENGTH: 222

**TYPE: pr
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APPLICANT: Huang, Shi
TITLE OF INVENTION: POLYPEPtides, Antibodies and Methods of Use FILE REFERENCE: P-LJ 5301
CURRENT APPLICATION NUMBER: US/10/200,012
CURRENT FILING DATE: 2002-07-18
PRIOR PLING DATE: 2001-07-18
PRIOR PLING DATE: 2001-07-18
NUMBER OF SEQ ID NOS: 46
SOFTWARE: FastSEQ for Windows Version 4.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1: || | | | | | : : || 36 QMPQPHPHQPNYPPQHSGSQVVVDRRELNERLDRLEKLLKEAEQRMNRQER 87
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Best Local Similarity 32.7%; Pred. No. 11;
Matches 17; Conservative 7; Mismatches
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53.8%; Pred. No. 19
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US-09-248-796A-21807
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Best Local Similarity 53.8
Matches 14; Conservative
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16 LPKEMTPVEPAAFAAELISRLEKLKLELE 44
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Job time : 5.51187 secs
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Best Local Similarity
Matches 12; Conserva
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APPLICANT: VENTER, J. Craig et al.

APPLICANT: VENTER, J. Craig et al.

TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED

TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF

FILE REFERENCE: CLOO1307

CURRENT APPLICATION NUMBER: uS/09/949,016

CURRENT FILING DATE: 2000-04-14

PRIOR PILING DATE: 2000-10-20

PRIOR PILING DATE: 2000-10-20

PRIOR FILING DATE: 2000-10-03

PRIOR FILING DATE: 2000-10-09

PRIOR FILING DATE: 2000-10-09

PRIOR FILING DATE: 2000-09-08
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APPLICANT: Nexi, Bruce P.
TITLE OF INVENTION: Enzymes for the Detection of Nucleic Acid Sequences FILE REFERENCE: PORS 04931
CURRENT APPLICATION NUMBER: US/09/758,282B
CURRENT FILING DATE: 2001-01-11
PRIOR APPLICATION NUMBER: 09/577,304
PRIOR FILING DATE: 2000-05-24
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         11 PRTHRLPKEMTPVEP----AAFAAELISRLEKLKLELE-SRHSLEERLQ 54
                                                         Length 1061;
                                                                                                                                                                           Score 57; DB 2; Length 392;
Pred. No. 43;
9; Mismatches 14; Indels
                                                           21.5%; Score 58.5; DB 2; Length 1 34.0%; Pred. No. 90; ive 10; Mismatches 16; Indels
                                                                                                                                                4 QVSLPHFPRTHRL---PKEMTPVEP--AAFAAELISRLEKLKLELES
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 6078
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Patent No. 6812339
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Kaiser, Michael W.
Lyamicheva, Natalie E.
Allawi, Hatim T.
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                                                         Query Match
Best Local Similarity 34.0*
Matches 16; Conservative
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SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 251
LENGTH: 832
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Best Local Similarity 36.7
Matches 18; Conservative
; ORGANISM: Human
US-10-200-012-4
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US-09-949-016-6078
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10; Indels

7; Mismatches

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CURRENT APPLICATION NUMBER: US/09/758,282B
CURRENT FILING DATE: 2001-01-11
PRIOR PLING DATE: 2001-01-11
PRIOR FILING DATE: 2000-05-24
NUMBER OF SEQ ID NOS: 280
SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence: Synthetic
US-09-758-2828-268
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Length 832;
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21.0%; Score 57; DB 2;
41.4%; Pred. No. 1e+02;
tive 7; Mismatches 10
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Pred. No. 1e+02;
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                                                                                                             216 LLKHLEQVKPASVREKILSHMEDLKLSLE
                                                                                     16 LPKEMTPVEPAAFAAELISRLEKLKLELE
                                                                                                                                                                                                                             ; Sequence 268, Application US/09758282B; Patent No. 6635463
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Kaiser, Michael W.
Lyamicheva, Natalie E.
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ORGANISM: Artificial Sequence
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Neri, Bruce P.
Ouery Match
Best Local Similarity 41.4'
Matches 12; Conservative
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us-09-587-574-3.rapbm

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                                                                         April 20, 2006, 15:57:53 ; Search time 18.7599 Seconds (without alignments) 1202.714 Million cell updates/sec
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272
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| cgn2 6/ptcdata/1/pubpaa/USO7 PUBCOMB.pep:*
2: /cgn2 6/ptcdata/1/pubpaa/USO8 PUBCOMB.pep:*
3: /cgn2 6/ptcdata/1/pubpaa/USO8 PUBCOMB.pep:*
4: /cgn2 6/ptcdata/1/pubpaa/USOB_PUBCOMB.pep:*
5: /cgn2 6/ptcdata/1/pubpaa/USIOB_PUBCOMB.pep:*
5: /cgn2 6/ptcdata/1/pubpaa/USIOB_PUBCOMB.pep:*
GenCore version 5.1.7
Copyright (c) 1993 - 2006 Biocceleration Ltd.
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-09-798-831-5
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                                                  OM protein - protein search, using sw model
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Maximum DB seq length: 200000000
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Match Length DB
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                                                                                                                                              Perfect score:
                                                                                                                                                                                   Scoring table:
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141.5
141.5
141.5
141.5
141.5
136.5
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60.5
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59.5
59.5
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                                                                                                                                                          Sequence:
                                                                                                                                                                                                                          Searched:
                                                                            Run on:
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Seguence 1342, Ap	Seguence 407, App	Seguence 108781,	Sequence 245, App	Sequence 140936,	Sequence 15, Appl	Sequence 16, Appl	Sequence 19, Appl	Sequence 20, Appl	Sequence 23, Appl	Seguence 24, Appl	Sequence 27, Appl	Sequence 28, Appl	Seguence 123393,	Sequence 42, Appl	Sequence 40, Appl	Seguence 52, Appl	Sequence 51, Appl	
US-10-732-923-1342	US-10-408-765A-407	US-10-437-963-108781	US-10-128-558-245	US-10-437-963-140936	US-10-799-369-15	US-10-799-369-16	US-10-799-369-19	US-10-799-369-20	US-10-799-369-23	US-10-799-369-24	US-10-799-369-27	US-10-799-369-28	US-10-437-963-123393	US-10-038-854-42	US-10-038-854-40	US-10-042-865-52	US-10-029-020-51	,
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369	410	415	459	614	833	833	833	833	833	833	833	833	931	2613	2628	2715	2715	
21.9	21.9	21.9	21.9	21.9	21.9	21.9	21.9	21.9	21.9	21.9	21.9	21.9	21.9	21.9	21.9	21.9	21.9	
59.5	59.5	59.5	59.5	59.5	59.5	59.5	59.5	59.5	59.5	59.5	59.5	59.5	59.5	59.5	59.5	59.5	59.5	
28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43	44	45	

## ALIGNMENTS

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Squence 1797, Application US/10723860

Squence 1797, Application US/10723860

Publication No. US20040253606A1

GRENEAL INPORMATION:
APPLICANT: Aziz, Natasha

APPLICANT: Zlotnik, Albert

TITLE OF INVENTION: Methods for Screening for Soft Tissue Sarcoma Modulators

TITLE OF INVENTION: Methods for Screening for Soft Tissue Sarcoma Modulators

FILE REFERENCE: 05882.0193.NPUS01

CURRENT APPLICATION NUMBER: US/10/723,860

CURRENT FILING DATE: 2003-11-26

PRIOR APPLICATION NUMBER: 60/429,739

PRIOR FILING DATE: 2002-11-26

NUMBER OF SEQ ID NOS: 8393

SGQ ID NOS: 8393

SGQ TU NO 1797
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 116, Application US/10751736
Publication No. US20040265230A1
GRNERAL INFORMATION:
GRNERAL INFORMATION:
GRNERAL INFORMATION:
APPLICANT: Wartinez, Robert
APPLICANT: Brown, Eugene
APPLICANT: Liv, Wei
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR DIAGNOSING AND TREATING COLON
TITLE OF INVENTION: CANCERS
FILE REFERENCE: AM100927 (031896-002000)
CURRENT FELING DATE: 2003-01-06
PRIOR APPLICATION NUMBER: US 105/10/751,736
CURRENT FILING DATE: 2003-01-06
PRIOR PILING DATE: 2003-01-06
PRIOR FILING DATE: 2003-01-06
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: PatentIn version 3.2
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Matches 53; Conservative
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RESULT 1
US-10-723-860-1797
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SEQ ID NO 116

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APPLICANT: Anisowicz, Anthony
APPLICANT: Bhat, Bheem
APPLICANT: Bhat, Bheem
APPLICANT: Danagora, Veronique
APPLICANT: Nobinson, John
APPLICANT: Yaworsky, Paul
TITLE OF INVENTION: Reagents and Method for Modulating DKK-Mediated Interactions
FILE REFERENCE: 032796-143
CURRENT APPLICATION NUMBER: US/10/182,936A
CURRENT FILING DATE: 2002-08-02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 900;
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                                                                                                                          TITLE OF INVENTION: THE HIGH BONE MASS GENE OF 11q13.3
TITLE OF INVENTION: THE HIGH BONE MASS GENE OF 11q13.3
FILE REFERENCE: 032796-021
FULE REFERENCE: 032796-021
CURRENT APPLICATION NUMBER: US/10/374,979
CURRENT FILING DATE: 2003-03-04
PRIOR FILING DATE: 2000-04-05
PRIOR APPLICATION NUMBER: US 09/543,771
PRIOR FILING DATE: 2000-04-05
PRIOR PELING DATE: 1999-01-13
PRIOR FILING DATE: 1999-01-13
PRIOR FILING DATE: 1998-01-13
PRIOR FILING DATE: 1998-10-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 4;
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Pred. No. 6e-08;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR FILING DATE: 2002-05-17
PRIOR APPLICATION NUMBER: US 60/291,311
PRIOR PILING DATE: 2001-05-17
PRIOR PILING DATE: 2002-02-01
PRIOR PILING DATE: 2002-02-01
PRIOR PILING DATE: 2002-03-04
PRIOR APPLICATION NUMBER: US 60/361,293
PRIOR PILING DATE: 2002-03-04
NUMBER OF SEQ ID NOS: 216
SUPTWARE: PRECSEQ for Windows Version 4.0
SEQ ID NO 91
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION NUMBER: PCT/US02/15982
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; sequence 91, Application US/10182936A; Publication No. US20040038860A1; GENERAL INFORMATION:
                                                                 ; Sequence 91, Application US/10374979; Publication No. US20030219793A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       52.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity 56.6
Matches 30; Conservative
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Best Local Similarity 56.6
Matches 30; Conservative
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; ORGANISM: Homo sapiens
US-10-374-979-91
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APPLICANT: Liu, Wei
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR DIAGNOSING AND TREATING AUTOIMMUNE
TITLE OF INVENTION: DISEASES
FILE REFERENCE: 031896-023000 (AM101331L)
CURRENT APPLICATION NUMBER: uS/10/786,720
CURRENT PILLNG DATE: 2004-02-26
NUMBER OF SEQ ID NOS: 21135
SOFTWARE: Patentin version 3.2
SEQ ID NO 36
LENGTH: 826
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APPLICANT: Wyeth
APPLICANT: U, Wa.
APPLICANT: O'Toole, Margot
APPLICANT: Liu, Will
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR DIAGNOSING AND TREATING AUTOIMMUNE
TITLE OF INVENTION: DISEASES
FILE REPERENCE: 031896-023000 (AM101331L)
CURRENT APPLICATION NUMBER: US/10/786,720
CURRENT FILING DATE: 2004-02-26
NUMBER OF SEQ ID NOS: 21135
SOFTWARE: Patentin version 3.2
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                                                                                                                Score 268; DB 5; Length 843;
Pred. No. 4.1e-23;
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                                                                                                                                                                   1; Indels
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Pred. No. 5.5e-08;
8; Mismatches 14
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; Sequence 35, Application US/10786720
; Publication No. US20040191818A1
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                            S-10-786-720-36
Sequence 36, Application US/10786720
Publication No. US20040191818A1
GENERAL INFORMATION:
                                                                                                                   Query Match
Best Local Similarity 98.1%;
Matches 53; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity 56.6%;
Matches 30; Conservative
LENGTH: 843
TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: PRT
CORGANISM: Homo sapiens
US-10-786-720-36
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ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Wyeth
                                                                        US-10-751-736-116
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LENGTH: 862
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Matches
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APPLICANT:
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Publication No. US20040221326A1
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Babij, Philip
APPLICANT: Bex, Frederick J. III
APPLICANT: Bex, Frederick J. III
APPLICANT: Bex, Frederick J. III
APPLICANT: Bex, Frederick J. III
APPLICANT: Bex, Frederick J. III
APPLICANT: Bodine, Peter Van Nest
IITLE OF INVENTION: Transgenic Animal Model of Bone Mass Modulation
FILE REFERENCE: 032796-212
24
NGOVSLPHFPRTHRLPKEMTPVEPAAFAAELISRLEKLKLELESRHSLEERLQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2 NGOVSLPHFPRTHRLPKEMTPVEPAAFAAELISRLEKLKLELESRHSLEERLO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 900;
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Pred. No. 6e-08;
8; Mismatches
                                                                                                                                                                                                                             FILE TREATORY NOWBER: US/10/477,238A CURRENT APPLICATION NUMBER: US/10/477,238A CURRENT FILING DATE: 2003-11-10
PRIOR FILING DATE: 2001-05-11
PRIOR FILING DATE: 2001-05-17
PRIOR FILING DATE: 2001-05-17
PRIOR PLING DATE: 2001-05-17
PRIOR PLING DATE: 2002-02-01
PRIOR APPLICATION NUMBER: US 60/353,058
PRIOR FILING DATE: 2002-03-04
NUMBER OF SEQ ID NOS: 812
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 670
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity 56.6%;
Matches 30; Conservative 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                             GRGANISM: Homo sapiens US-10-477-238A-670
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: Homo sapiens
                                                                                     10-477-238A-670
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LENGTH: 900
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CURRENT APPLICATION NUMBER: US/10/477,173
CURRENT FILING DATE: 2003-11-0
PRIOR PLING DATE: 2001-55-11
PRIOR APPLICATION NUMBER: US 60/290,071
PRIOR APPLICATION NUMBER: US 60/291,311
PRIOR PILING DATE: 2001-05-17
PRIOR FILING DATE: 2001-05-17
PRIOR FILING DATE: 2002-02-01
PRIOR APPLICATION NUMBER: US 60/351,293
PRIOR PLING DATE: 2002-02-01
PRIOR PLING DATE: 2002-02-01
                                              Gaps
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                                                                                                                      2 NGOVSLPHFPRTHRLPKEMTPVEPAAFAAELISRLEKLKLELESRHSLEERLQ 54
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    Length 900;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 900;
Query Match 52.0%; Score 141.5; DB 5; Length 9 Best Local Similarity 56.6%; Pred. No. 6e-08; Matches 30; Conservative 8; Mismatches 14; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          14; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 141.5; DB 5;
Pred. No. 6e-08;
8; Mismatches 14;
                                                                                                                                                                                                                                                                                                                      APPLICANT: Genome Therapeutics Corporation and APPLICANT: Allen, Kristina M. APPLICANT: Yaworsky, Paul APPLICANT: Morales, Arturo J. APPLICANT: Anisowicz, Anthony APPLICANT: Liu, Wei
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQ ID NOS: 1086
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 270, Application US/10092900A Publication No. US20040043382A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                         ; Sequence 670, Application US/10477173; Publication No. US20050070699A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Padigaru, Muralidhara
APPLICANT: Spytek, Kimberly A.
APPLICANT: Shenoy, Suresh G.
APPLICANT: Taupier Jr., Raymond J.
APPLICANT: Pena, Carol E.A.
APPLICANT: Li, Li
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity 56.6%;
Matches 30; Conservative
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Gangolli, Esha A.
Vernet, Corine A.M
Guo, Xiaojia Sasha
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Zerhusen, Bryan D.
Gusev, Vladimir Y.
Ji, Weizhen
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Miller, Charles E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Kekuda, Ramesh
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 10
US-10-092-900A-270
                                                                                                                                                                                                                                       US-10-477-173-670
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LENGTH: 900
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APPLICANT: KLEIN, Peter S.

TITLE OF INVENTION: AXIN DOMAIN-LIKE POLYPEPTIDE INHIBITORS OF GLYCOGEN
TITLE OF INVENTION: SYNTHASE KINASE 3 (BETA) ACTIVITY AND ACTIVATORS OF WNT
TITLE OF INVENTION: SIGNALING
FILE REPERENCE: 2095-66.0391/306U1
CURRENT APPLICATION NUMBER: US/09/798,831
CURRENT FILING DATE: 2001-03-01
RIOR APPLICATION NUMBER: US 60/186,141
PRIOR APPLICATION NUMBER: US 60/186,141
PRIOR APPLICATION NUMBER: US 60/186,141
SEQ ID NOS: 12
SOFTWARE: PatentIN Ver. 2.1
SEQ ID NO 6 .
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TITLE OF INVENTION: AXIN DOMAIN-LIKE POLYPEPTIDE INHIBITORS OF GLYCOGEN
TITLE OF INVENTION: SYNTAKE KINASE 3 (BETA) ACTIVITY AND ACTIVATORS OF WNT
TITLE OF INVENTION: SYNTAKES X (BETA) ACTIVITY AND ACTIVATORS OF WNT
TITLE OF INVENTION: SYNTAKEN (BETA)
FILE REFERENCE: 209596.0391/306U1
CURRENT APPLICATION NUMBER: US 60/186,141
PRIOR APPLICATION NUMBER: US 60/186,141
PRIOR APPLICATION NUMBER: US 60/186,141
SAPTOR FILING DATE: 2000-03-01
NUMBER OF SEQ ID NOS: 12
SOFTWARE: PAECHLIN Ver. 2.1
                                                                                                                                                     Gaps
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                                                                                                                                                                                                                                          54
                                                                                                                                                                                                        1 ANGQVSLPHFPRTHRLPKEMTPVEPAAFAAELISRLEKLKLELESRHSLEERLQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CTHER INFORMATION: Description of Artificial Sequence: Rat axil; OTHER INFORMATION: residues 362-386
US-09-798-831-6
                                                                                                  DB 3; Length 842;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CTHER INFORMATION: Description of Artificial Sequence: Murine COTHER INFORMATION: conductin residues 362-386
US-09-798-831-7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      43.4%; Score 118; DB 3; Length 25; 100.0%; Pred. No. 5.4e-07; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          43.4%; Score 118; DB 3; Length 25.
                                                                                               ; Score 136.5; DB 3
; Pred. No. 2.2e-07;
10; Mismatches 14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    20 MTPVEPAAPAAELISRLEKLKLELE 44
                                                                                                                                                                                                                                                                                                                                                                                            Sequence 6, Application US/09798831
Patent No. US20010052137A1
GENERAL INFORMATION:
APPLICANT: KLEIN, Peter S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 7, Application US/09798831
Patent No. US20010052137A1
                                                                                                  Query Match
Best Local Similarity 53.7%;
Matches 29; Conservative 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity 100.0
Best Local Similarity
Section 25, Conservative
; TYPE: PRT
; ORGANISM: Xenopus laevis
US-09-798-831-8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: KLEIN, PELER S.

TITLE OF INVENTION: AXIN DOMAIN-LIKE POLYPEPTIDE INHIBITORS OF GLYCOGEN
TITLE OF INVENTION: SYNTHESE KINASE 3 (BETA) ACTIVITY AND ACTIVATORS OF WNT
TITLE OF INVENTION: SIGNALING
FILE REPERENCE: 209596.0391/30601
CURRENT APPLICATION NUMBER: US/09/798,831
CURRENT FILING DATE: 2001-03-01
PRIOR APPLICATION NUMBER: US 60/186,141
PRIOR FILING DATE: 2000-03-01
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 768
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2 NGOVSLPHFPRTHRLPKEMTPVEPAAFAAELISRLEKLKLELESRHSLEERLO 54
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Pred. No. 6.1e-08;
8; Mismatches 14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: USSN 60/274,322
PRIOR PLILING DATE: 2002-03-07
PRIOR PLILING DATE: 2001-03-08
PRIOR PELLING DATE: 2001-03-08
PRIOR PELLING DATE: 2001-03-08
PRIOR PELLING DATE: 2001-04-13
PRIOR PELLING DATE: 2001-04-13
PRIOR PELLING DATE: 2001-04-08
PRIOR PELLING DATE: 2001-03-08
PRIOR PELLING DATE: 2001-03-08
PRIOR PELLING DATE: 2001-03-08
PRIOR PELLING DATE: 2001-03-08
PRIOR PELLING DATE: 2001-03-08
PRIOR PELLING DATE: 2001-03-08
PRIOR PELLING DATE: 2001-03-08
PRIOR PELLING DATE: 2001-03-08
PRIOR PELLING DATE: 2001-03-27
PRIOR PELLING DATE: 2001-03-27
PRIOR PELLING DATE: 2001-03-27
PRIOR PELLING DATE: 2001-03-38
PRIOR PELLING DATE: 2001-03-39
PRIOR PELLING DATE: 2001-03-30
                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/10/092,900A CURRENT FILING DATE: 2002-03-07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION NUMBER: USSN 60/294,899
PRIOR FILING DATE: 2001-05-31
PRIOR APPLICATION NUMBER: USSN 60/287,424
PRIOR FILING DATE: 2001-04-30
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Patent No. US20010052137A1
                                                                                                                                                                                                                                                                                                                                                           Burgess, Catherine E
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Catterton, Elina
Leite, Mario W.
                                                        Uriel M
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Best Local Similarity 56.6%;
Matches 30; Conservative
     Fernandes, Elma R.
                                                                                                                                                                                                                                                                   Alsobrook, John P.
Lepley, Denise M.
                                                                                                                                    Anderson, David W.
                               Casman, Stacie J.
Malyankar, Uriel
                                                                                                                                                                                                                                               Haihong
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US-10-092-900A-270
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TITLE OF INVENTION: AXIN DOMAIN-LIKE POLYPEPTIDE INHIBITORS OF GLYCOGEN
TITLE OF INVENTION: SYNTHASE KINASE 3(BETA) ACTIVITY AND ACTIVATORS OF WNT
TITLE OF INVENTION: SIGNALING
FILE REFERENCE: 209596.0391/306U1
CURRENT APPLICATION NUMBER: US/09/798,831
PRIOR APPLICATION NUMBER: US 60/186,141
PRIOR APPLICATION NUMBER: US 60/186,141
PRIOR PILING DATE: 2000-03-01
WUMBER OF SEC ID NOS: 12
SOFTWARE: PATENTIN VET. 2.1
                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OTHER INFORMATION: Description of Artificial Sequence: Human axin 2 OTHER INFORMATION: residues 362-386
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 41.9%; Score 114; DB 3; Length 25; Best Local Similarity 96.0%; Pred. No. 1.6e-06; Matches 24; Conservative 0; Mismatches 1; Indels
                   Indels
Pred. No. 5.4e-07;
Mismatches 0;
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                                                                                20 MTPVEPAAFAAELISRLEKLKLELE 44
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                                                                                                                                                                                                   Sequence 5, Application US/09798831
Patent No. US20010052137A1
GENERAL INFORMATION:
APPLICANT: KLEIN, Peter S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gerlach, Valerie
Taupier Jr, Raymond J.
Gusev, Vladimir Y.
Colman, Steven D.
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Padigaru, Muralidhara
Best Local Similarity 100.0%; P
Matches 25; Conservative 0;
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Grosse, William M.
Alsobrook II, John P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Burgess, Catherine E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: PRT ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Anderson, David W.
Rastelli, Luca
Miller, Charles E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Zerhusen, Bryan
Patturajan, Meera
Shimkets, Richard
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Rieger, Daniel K.
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US-10-072-012-490
                                                                                                                                                                                    US-09-798-831-5
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LENGTH: 25
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14078, A
2, Appli
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Sequence 17616, A
Sequence 23144, A
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                                                                  April 20, 2006, 16:00:23; Search time 2.99208 Seconds (without alignments) 794.148 Million cell updates/sec
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Description
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Sequence 8
Sequence 4
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Sequence
Sequence
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Sequence
Sequence
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272
1 ANGQVSLPHFPRTHRLPKEM.....RLEKLKLELESRHSLEERLQ
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1: /SIDSS/ptodata/2/pubpaa/USO8 NEW_PUB.pep:*
2: /SIDSS/ptodata/2/pubpaa/USO6_NEW_PUB.pep:*
3: /SIDSS/ptodata/2/pubpaa/USO7_NEW_PUB.pep:*
4: /SIDSS/ptodata/2/pubpaa/USO7_NEW_PUB.pep:*
5: /SIDSS/ptodata/2/pubpaa/USIO_NEW_PUB.pep:*
6: /SIDSS/ptodata/2/pubpaa/USIO_NEW_PUB.pep:*
7: /SIDSS/ptodata/2/pubpaa/USIO_NEW_PUB.pep:*
8: /SIDSS/ptodata/2/pubpaa/USIO_NEW_PUB.pep:*
        GenCore version 5.1.7
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US-11-096-568A-23143
US-11-096-051-2
US-11-113-424-51
US-11-096-051-10
US-11-096-051-8
US-11-121-438-4
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US-11-188-298-17616
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US-11-079-463-5959
US-11-087-099-6670
US-10-467-657-7122
US-10-481-935A-208
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                                                                                                                                                                                                             of hits satisfying chosen parameters:
                                                                                                                                                                                         225428 segs, 44002918 residues
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Maximum Match 100%
Listing first 45 summaries
                                                 - protein search, using sw model
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Gapop 10.0 , Gapext 0.5
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Maximum DB seq length: 2000000000
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Query
Match Length DB
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Perfect
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Sequence 215, Application US/10501035
Publication No. US20060046249A1
GENERAL INFORMATION:
GENERAL INFORMATION:
GENERAL INFORMATION:
TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
ACTIVITY OF COMPOUNDS THAT INTERACT WITH PROTEIN TYROSINE KINA:
TITLE REPERENCE:
D0185 PCT
CURRENT APPLICATION NUMBER:
US/10/501,035
CURRENT FILING DATE:
2004-07-09
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                                                                                          1173, Ap
4, Appli
2, Appli
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2, Appli
6, Appli
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                                                                                               Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequence In the sequ
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; Publication No. US20060075522A1
; GENERAL INFORMATION:
; APPLICANT: Abad, Mark S. et al.
; TITLE OF INVENTION: GENES AND USES FOR PLANT IMPROVEMENT
; FILE REFERENCE: 38-21(53452)B
; CURRENT APPLICATION NUMBER: 08/11/188,298
; CURRENT FILING DATE: 2005-07-22
; PRIOR APPLICATION NUMBER: 60/592,978
; NUMBER OF SEQ ID NOS: 22569
; SEQ ID NO 16414
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        52.0%; Score 141.5; DB 6; ilarity 56.6%; Pred. No. 3.4e-09; Conservative 8; Mismatches 14;
                     US-11-087-099-9570
US-11-087-099-10782
US-11-087-099-1173
US-11-046-668-2
US-11-046-668-2
US-10-055-877-36
US-10-055-877-36
US-11-087-099-1058
US-11-087-099-1058
US-11-118-122-2
US-10-453-195-6
US-11-169-041-183
US-10-131-826A-270
US-10-131-826A-270
US-10-131-826A-270
US-10-131-826A-270
US-10-131-826A-270
US-10-131-826A-270
US-10-821-234-1185
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PRIOR APPLICATION NUMBER: US 60/350,061
PRIOR FILING DATE: 2002-01-18
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SOFTWARE: Patentin version 3.2
SEQ ID NO 215
LENGTH: 900
487
1758
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267
2267
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329
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ORGANISM: Homo sapiens
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ORGANISM: Oryza sativa
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Best Local Similarity
Them 30; Conserv?
  US-11-188-298-16414
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Sequence Sequence Sequence

Sequence Sequence

US-11-096-568A-8496 US-11-096-568A-6619 US-11-096-568A-6618

US-11-096-568A-6620 US-11-096-568A-8497

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APPLICANT: Alexandrov, Nickolai et al.
TITLE OF INVENTION: Sequence-Determined DNA Fragments and Corresponding Polypeptide
TITLE OF INVENTION: Therby
TITLE OF INVENTION: Therby
FILE REPERENCE: 2750-1592PUS2
CURRENT APPLICATION NUMBER: US/11/096,568A
CURRENT FILING DATE: 2005-04-01
NUMBER OF SEQ ID NOS: 34471
SEQ ID NO 21843
LENGTH: 431
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 23144, Application US/11096568A

Publication No. US20060048240A1

GENERAL INFORMATION:
APPLICAMY: Alexandrow, Nickolai et al.
TITLE OF INVENTION: Sequence-Determined DNA Fragments and Corresponding Polypeptide
TITLE OF INVENTION: Therby
FILE REFERENCE: 2750-1592PUS2
CURRENT APPLICATION WHOBER: US/11/096,568A
CURRENT RILING DATE: 2005-04-01
NUMBER OF SEQ ID NOS: 34471
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                                 APPLICANT: Abad, Mark S. et al.
TITLE OF INVENTION: GENES AND USES FOR PLANT IMPROVEMENT
FILE REPRENCE: 38-21(53452)B
CURRENT APPLICATION NUMBER: US/11/188,298
CURRENT FILING DATE: 2005-07-22
PRIOR APPLICATION NUMBER: 60/592,978
PRIOR PELING DATE: 2004-07-31
NUMBER OF SEQ ID NOS: 22569
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; OTHER INFORMATION: Ceres Seq. ID no. 12411437
US-11-096-568A-23144
                                                                                                                                                                                                                                                                                                                                   Score 59.5; D
Pred. No. 11;
7; Mismatches
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Pred. No. 12;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: PRT
ORGANISM: Zea mays subsp. mays
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34.5%;
US20060075522A1
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Best Local Similarity 35.8%;
Matches 19; Conservative
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Best Local Similarity 34.5
Matches 19; Conservative
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US-11-188-298-17616
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US-11-096-568A-23143
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LENGTH: 381
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TITLE OF INVENTION: Sequence-Determined DNA Fragments and Corresponding Polypeptides
TITLE OF INVENTION: Therby
FILE REFERENCE: 2750-1592PUS2
CURRENT APPLICATION NUMBER: US/11/096,568A
CURRENT FILING BATE: 2005-04-01
NUMBER OF SEQ ID NOS: 34471
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                                     Length 213;
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                                                                                                                                                    PRDDGQPPDLAGMDPAALQAELLRRHAGPSFGFVKLRLGVRRSQEE 103
                                                                                                                     11 PRTHRLPKEMTPVEPAAFAAELISR----LEKLKLELESRHSLEE 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          11 PRTHRLPKEMTPVEPAAFAAELISR----LEKLKLELESRHSLEE 51
                                                                               Indels
                                                                                                                                                                                                                                                                                                                                   APPLICANT: Abad, Mark S. et al.
TITLE OF INVENTION: GENES AND USES FOR PLANT IMPROVEMENT
FILE REFERENCE: 38-21(53452)8
CURRENT APPLICATION NUMBER: 2005-07-22
PRIOR APPLICATION NUMBER: 60/592,978
PRIOR FILING DATE: 2004-07-31
NUMBER OF SEQ ID NOS: 22569
SEQ ID NO 18243
LENGTH: 213
                                                                               18;
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                                       DB 7;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; ORGANISM: Oryza sativa (japonica cultivar-group)
US-11-188-298-18243
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| LOCATION: (1)..(323)
| OTHER INFORMATION: Ceres Seq. ID no. 12411438
| US-11-096-568A-23145
                                       Score 59.5; DE
Pred. No. 5.8;
                                                                               6; Mismatches
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                                                                                                                                                                                                                                                                        Sequence 18243, Application US/11188298
Publication No. US20060075522A1
GENERAL INFORMATION:
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US-11-188-298-17616
; Sequence 17616, Application US/11188298
                                       21.9%;
37.0%;
                               Query Match
Best Local Similarity 37.0°
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Best Local Similarity 37.0.
These 17; Conservative
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Matches 19, Conserv
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-11-188-298-18243 (
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  US-11-188~298-16414
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LENGTH: 323
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APPLICANT: Kekuda, Ramesh
APPLICANT: Kekuda, Ramesh
APPLICANT: Kekuda, Ramesh
APPLICANT: MacLachlan, Timothy K
APPLICANT: MacLachlan, Timothy K
APPLICANT: Rastelli, Luca
APPLICANT: Vernet, Corine
APPLICANT: Vernet, Corine
APPLICANT: Vernet, Corine
TITLE OF INVENTION: Tar-M3 Polypeptides and Polynucleotides and their Methods of Us-
TITLE REPERBNCE: Attorney Docket No. Cura 967
CURRENT APPLICATION NUMBER: US/11/096,051
PRIOR APPLICATION NUMBER: 10/038,854
PRIOR APPLICATION NUMBER: 10/038,854
PRIOR APPLICATION NUMBER: 10/455,772
PRIOR APPLICATION NUMBER: 60/557,978
PRIOR APPLICATION NUMBER: 60/557,978
PRIOR APPLICATION DATE: 2003-06-04
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Pred. No. 1.2e+02;
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Pred. No. 1.2e+02;
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PRIOR APPLICATION NUMBER: 60/315,61,
PRIOR FILING DATE: 2001-08-29
PRIOR PLING DATE: 2001-07-24
PRIOR PLING DATE: 2001-07-24
PRIOR APPLICATION NUMBER: 60/322,358
PRIOR APPLICATION NUMBER: 60/294,075
PRIOR APPLICATION NUMBER: 60/294,075
PRIOR FILING DATE: 2001-09-14
PRIOR PILING DATE: 2001-05-29
PRIOR FILING DATE: 2001-05-29
PRIOR FILING DATE: 2001-05-29
PRIOR FILING DATE: 2001-05-62
SOFTWARE PLANG DATE: 2001-05-62
SOFTWARE PARENTE: PARENTE: 2001-05-63
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQ ID NOS: 38
SOFTWARE: CuraSeqList version 0.1
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Best Local Similarity 26.3
Matches 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    20; Conservative
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; ORGANISM: Homo sapiens
US-11-096-051-10
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APPLICANT: MacLachlan, Timothy K
APPLICANT: MatLachlan, Timothy K
APPLICANT: Rastelli, Luca
APPLICANT: Restelli, Luca
APPLICANT: Vernet, Corine
APPLICANT: Ettenberg, Seth
TITLE OF INVENTION: Ten-M3 Polypeptides and Polynucleotides and their Methods of Use
FILE REFERENCE: Attorney Docket No. Cura 967
CURRENT APPLICATION NUMBER: US/11/096,051
CURRENT FILING DATE: 2005-03-30
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; Bequence 51, Application No. US20050260713A1
; GENERAL INFORMATION:
; APPLICANT: Gangolli et al.
; TITLE OF INVENTION: Polypeptides and Nucleic Acids Encoding Same; FILE REFERENCE: 21402-225
; CURRENT FILING DATE: 2005-04-21
; PRIOR FILING DATE: 2000-04-21
; PRIOR FILING DATE: 2000-12-19
; PRIOR FILING DATE: 2010-108-10
; PRIOR PLICATION NUMBER: 60/257,314
; PRIOR FLING DATE: 2010-08-10
; PRIOR FLING DATE: 2010-12-0
; PRIOR FLING DATE: 2010-12-0
; PRIOR FLING DATE: 2010-12-0
; PRIOR FLING DATE: 2010-108-10
; PRIOR FLING DATE: 2010-108-10
                                                                                                                                                                                  Length 431;
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Pred. No. 1.2e+02;
9; Mismatches 18;
             ; NAME/KEY: misc_feature

; LOCATION: (1)...(431)

; OTHER INFORMATION: Ceres Seq. ID no. 12411436

US-11-096-568A-23143
                                                                                                                                                                           Query Match
21.9%; Score 59.5; DE
Best Local Similarity 34.5%; Pred. No. 14;
Matches 19; Conservative 11; Mismatches
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PRIOR APPLICATION NUMBER: 10/038,854
PRIOR FILING DATE: 2001-12-31
PRIOR APPLICATION NUMBER: 10/455,772
PRIOR FILING DATE: 2003-06-04
PRIOR FILING DATE: 2004-03-30
NUMBER OF SEQ ID NOS: 38
SOFTWARE: CuraSeqList version 0.1
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Publication No. US20050244868A1
GENERAL INFORMATION:
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26.3%;
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Best Local Similarity 26.3
Matches 20; Conservative
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ORGANISM: Homo sapiens
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LENGTH: 2715
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Sequence 5959, Application US/11079463
Publication No. US20060073161A1
GENERAL INFORMATION:
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO BACTEROIDES P.
TITLE OF INVENTION: POR DIAGNOSTICS AND THERAPEUTICS
TITLE OF INVENTION: VOMBER: US/11/079,463
CURRENT APPLICATION NUMBER: US 60/128,705
PRIOR PEDILICATION NUMBER: US 60/128,705
PRIOR FILING DATE: 1999-04-09
PRIOR FILING DATE: 1999-04-09
PRIOR PELICATION NUMBER: US 09/540,209
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Publication No. US20050265985A1
GENERAL INFORMATION:
APPLICANT: CHODOSH, Lewis A
APPLICANT: GARDNER, Heather P
TITLE OF INVENTION: HORWONALLY UP-REGULATED, NEU-TUMOR-ASSOCIATED KINASE
FILE REPERENCE: 22253-070821
CURRENT APPLICATION NUMBER: US/11/121,419
CURRENT FILING DATE: 2005-05-04
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                                                                                 APPLICANT: Abad, Mark S. et al.
TITLE OF INVENTION: GENES AND USES FOR PLANT IMPROVEMENT
FILE REFERENCE: 38-21(53452)B
CURRENT APPLICATION NUMBER: US/11/188,298
CURRENT FILING DATE: 2005-07-22
PRIOR APPLICATION NUMBER: 60/592,978
PRIOR PELLING DATE: 2004-07-31
NUMBER OF SEQ ID NOS: 22569
SEQ ID NO 14078
                                                                                                                                                                                                                                                                                                                                                                                                                          Score 57.5; DB 7;
Pred. No. 13;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION NUMBER: US/10/032,256
PRIOR PILING DATE: 2001-12-21
PRIOR PILING DATE: 2000-12-21
PRIOR FILING DATE: 2000-12-21
PRIOR FILING DATE: 2000-12-21
PRIOR FILING DATE: 2000-12-21
            Sequence 14078, Application US/11188298
Publication No. US20060075522A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                               ; TYPE: PRT
; ORGANISM: Aquifex aeolicus VF5
US-11-188-298-14078
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Best Local Similarity 28.6%;
Matches 16; Conservative
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SEQ ID NO 2
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US-11-121-419-2
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US-11-079-463-5959
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                                                                                   APPLICANT: Kekuda, Ramesh
APPLICANT: MacLachlan, Timothy K
APPLICANT: MacLachlan, Timothy K
APPLICANT: MacLachlan, Timothy K
APPLICANT: Rastelli, Luca
APPLICANT: Brenel, Corine
APPLICANT: Ettenberg, Seth
TITLE OF INVENTION: Ten-M3 Polypeptides and Polynucleotides and their Methods of Use
TITLE REFERENCE: Attorney Docket No. Cura 967
CURRENT APPLICATION NUMBER: US/11/096,051
CURRENT APPLICATION NUMBER: 10/038,854
PRIOR APPLICATION NUMBER: 10/038,854
PRIOR FILING DATE: 2001-12-31
PRIOR FILING DATE: 2003-06-04
PRIOR FILING DATE: 2003-06-04
PRIOR FILING DATE: 2003-06-04
PRIOR FILING DATE: 2003-06-03-30
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GENERAL INFORMATION:
TITLE OF INVENTION: Polypeptides, Antibodies and Methods of Use; FILLE REFERENCE: P-LJ 5301; CURRENT APPLICATION NUMBER: US/11/121,438; CURRENT FILING DATE: 2005-05-03; PRIOR APPLICATION NUMBER: US/10/200,012; PRIOR APPLICATION NUMBER: US/10/200,012; PRIOR PLING DATE: 2002-07-18; PRIOR PLING DATE: 2001-07-18
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34.0%; Pred. No. 53;
tive 10; Mismatches
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 4
LENGTH: 1061
               Sequence 8, Application US/11096051
Publication No. US20050244868A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQ ID NOS: 38
SOFTWARE: CuraSeqList version 0.1
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Matches 16; Conservative
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Best Local Similarity 26.3
Matches 20; Conservative
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LENGTH: 2725
US-11-096-051-8
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RESULT 13

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Search completed: April 20, 2006, 16:07:53 Job time : 3.99208 secs

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Sequence 9, Appli
Sequence 7, Appli
Sequence 7, Appli
Sequence 11597, A
Sequence 17434, A
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Sequence 2760, Ap
Sequence 2343, Ap
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Sequence 11347, A
Sequence 11650, A
                                                                                                April 20, 2006, 15:30:33 ; Search time 5.76517 Seconds (without alignments) 989.497 Million cell updates/sec
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                      1 QIREDEEKEGSEQALSSRDG......vLKTPGCQSPGVGRYSPRSR 69
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Sequence 2,
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GenCore version 5.1.7
Copyright (c) 1993 - 2006 Biocceleration Ltd.
                                                                                                                                                                                                                                                                                                                             572060
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /cgn2_6/ptodata/1/iaa/5_COMB.pep:*
/cgn2_6/ptodata/1/iaa/6_COMB.pep:*
/cgn2_6/ptodata/1/iaa/H_COMB.pep:*
/cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*
/cgn2_6/ptodata/1/iaa/RE_COMB.pep:*
/cgn2_6/ptodata/1/iaa/RE_COMB.pep:*
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-09-252-991A-17434
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-09-902-540-11650
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                                                                                                                                                                                                                                                                                                                            Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            .09-220-157A-4
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Maximum Match 100%
Listing first 45 summaries
                                                                  OM protein - protein search, using sw model
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                                                                                                                                                                                                                                                                                                                                                              Minimum DB seq length: 0
Maximum DB seq length: 2000000000
                                                                                                                                                                      US-09-587-574-4
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Match Length DB
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113.5
72.5
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68.5
67.5
66.5
65.5
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9, Appli	Sequence	US-08-476-900A-9	7	1582	16.4	59	45
9, Appli	Sequence	US-08-404-531B-9	Н	1582	16.4	59	44
29, Appl	Sequence	US-08-488-546A-29	~	1498	16.4	59	43
29, Appl	Sequence	US-08-476-900A-29	~	1498	16.4	59	42
29, Appl	Sequence	US-08-404-531B-29	ч	1498	16.4	59	41
12646, A	Sequence	US-09-489-039A-12646	7	837	16.4	59	40
17141, A	Sequence	US-09-252-991A-17141	7	485	16.4	59	39
24897, A	Sequence	US-09-252-991A-24897	~	360	16.4	59	38
4, Appli	Sequence	US-09-920-804-4	~	1088	16.5	59.5	37
17964, A	Sequence	US-09-252-991A-17964	7	260	16.8	60.5	36
2474, Ap	Sequence	US-10-104-047-2474	~	897	16.9	61	35
19940, A	Sequence	US-09-252-991A-19940	7	477	16.9	61	34
17831, A	Sequence	US-09-252-991A-17831	~	283	16.9	61	33
43189, A	Seguence	US-09-270-767-43189	7	313	17.1	61.5	32
58525, A	Sequence	US-09-270-767-58525	~	203	17.1	61.5	31
22041, A	Sequence	US-09-248-796A-22041	7	66	17.1	61.5	30
15777, A	Seguence	US-09-248-796A-15777	~	545	17.2	62	29
17, Appl	Sequence 17, Appl	US-10-332-795-17	~	409	17.2	62	28

## ALIGNMENTS

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RESULT 1

US-08-890-865A-10

US-08-890-865A-10

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US-08-890-865A-10

US-08-890-865A-10

US-08-890-865A-10

US-08-890-865A-10

US-08-890-865A-10

US-08-890-865A-10

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US-08-890-865A-10
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34 ----SYEEDPQTILDDHLSRVLKTPGCQSPGVGRYSPRSR 69

3 REDEEK------EGSEQALSSRDGAPVQHPLALLPSG--

Gaps

38;

Ouery Match 40.6%; Score 146; DB 2; Length 855; Best Local Similarity 34.7%; Pred. No. 1e-09; Matches 35; Conservative 14; Mismatches 14; Indels

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Sequence 2, Application US/09734674;
Patent No. 6498022;
GENERAL INFORMATION:
APPLICANT: WEL, Ming-Hui et al
TITLE OF INVENTION: ISOLATED HUMAN TRANSPORTER PROTEINS,
TITLE OF INVENTION: NUCLEIC ACID MOLECULES ENCODING HUMAN TRANSPORTER PROTEINS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      440 REAEBKLEERLKRVRMEEEGEDGDPSSGPPGPC-HKLPPAPAWHHFPPRLCWTWACAGLR 498
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 2, Application US/09920804

Patent No. 6673899

GENERAL INFORMATION:

APPLICANT: Seino, Susumu; JCR Pharmaceuticals Co., Ltd.

TITLE OF INVERTION: Sodium Ion-Driven Chloride/Bi-Carbonate Exchanger

FILE REFERENCE: GP4

CURRENT APPLICATION NUMBER: US/09/920,804

CURRENT FILING DATE: 2001-08-03

NUMBER OF SEQ ID NOS: 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       31;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 900;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         22; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           18; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 3 REDEEK-------EGSEQALSSRDGAPVQHPLALLPS--
                                                                                            SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/890,865A FILMED DATE: 10-JUL-1997 CLASSIFICATION: 435 CLASSIFICATION: 435 NAME: White, John P REGISTRATION NUMBER: 28,678
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 72.5; DB 2;
Pred. No. 3.4;
6; Mismatches 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  33 GSYEEDPQTILDDHLSRVLKTPGCQSPGVGRYSPRS 68
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      5e-05
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 113.5;
                                                                                                                                                                                                                                                                 REFERENCE/DOCKET NUMBER: 0575/54249
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212)278-0400
                                                           COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                31.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Best Local Similarity 38.3%;
Matches 18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                   : 900 amino acids
amino acid
                                                                                                                                                                                                                                                                                                                               TELEFAX: (212)391-0526
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity 34.49
Matches 33; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MOLECULE TYPE: Protein
ZIP: 10036
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                        STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: PRT
; ORGANISM: Mus musculus
US-09-920-804-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQ ID NO 2
LENGTH: 1088
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US-08-890-865A-4
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  473 MQMRDAHEENPESILDEHVQRVMKTPGCQSPGPGRHSPKPR 513
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/890,865A FILING DATE: 10-JUL-1997 CLASSIFICATION: 435 ATTONREY/AGENT INPOME/ATION:
NAME: White, John P
REGISTRATION NUMBER: 28,678
REGISTRATION NUMBER: 0575/54249
TELECOMMULCATION: INFORMATION:
TELECOMMULCATION: 1000RAMILATION:
                                                                                                                                                                                                                                AXIN GENE AND USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AXIN GENE AND USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
38.8%; Score 139.5; DB 2
Best Local Similarity 35.3%; Pred. No. 8.4e-09;
Matches 30; Conservative 14; Mismatches 20
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                                                                                                                                                                                                                                  STREET: 1185 Avenue of the Americas CITY: New York COUNTRY: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADDRESSEE: Cooper & Dunham LLP
STREET: 1185 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    44 DDHLSRVLKTPGCQSPGVGRYSPRS 68
                                                                                                                                                                  GENERAL INFORMATION:
APPLICANT: Constantini, Franklin
APPLICANT: Zeng, Li
TITLE OF INVENTION: AXIN GENE AND U
NUMBER OF SEQUENCES: 23
CORRESPONDENCE ADDRESS:
ADDRESSEE: COOPER & Dunham LLP
                                                                                                                                                                                                                                                                                                                                                                                                 ZIP: 10036
COMPUTER REARABLE FORM:
MEDIUM TYPE: Floppy disk
CMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
APPLICANT: Constantini, Franklin
APPLICANT: Zeng, Li
TITLE OF INVENTION: AXIN GENE AND
TITLE OF SEQUENCES: 23
CORRESPONDENCE ADDRESS:
                                                                                                                      Sequence 1, Application US/08890865A
Patent No. 6307019
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 4, Application US/08890865A
Patent No. 6307019
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          992 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEFAX: (212)391-0526
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 992 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TOPOLOGY: linear
MOLECULE TYPE: Protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      linear
                                                                                                       -08-890-865A-1
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                                                                                                                                                   Gaps
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                                                                                                         Length 856;
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                                                                                                                                                 12; Indels
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                                                                                                                                                                                                                                                                                                                               Sequence 2143, Application US/10104047

Patent No. 6943241

GENERAL INPORMATION:
FORESTAL INFORMATION:
FILE OF INVENTION:
FILE REFERENCE: H1-A0105

CURRENT APPLICATION NUMBER:
FRIGHE TAILOR DATE: 2002-03-25

PRIOR FILING DATE: 2002-03-25

NUMBER OF SEQ ID NOS: 4096

SOFTWARE: Patentin Ver. 2.1

LENGTH: 1034
                                                                                                                                                                                                                              94 LTDSDGNPLQAGSSGTHPLAIXPQGEWVSPEPTLIDGH 131
                                                                                                                                                                                           15 LSSRDGAPVO-----HPLALLPSGSYEEDPQTILDDH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; Score 69.5; DI
; Pred. No. 7.8;
15; Mismatches
                                                                                                                                                 6; Mismatches
                                                                                                       Score 70;
Pred. No. 5
                   TYPE: PRT
; ORGANISM: Corynebacterium glutamicum
US-09-605-703B-2760
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
APPLICANT: Scanlan, Thomas S
APPLICANT: Baxter, John D
APPLICANT: Fletterick, Robert J
APPLICANT: Wagner, Richard L
APPLICANT: Kushner, Peter J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 9, Application US/08764870
Patent No. 6236946
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                43 LDDHLSRV---LKTPGCQSP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity 28.8%;
Matches 23; Conservative 11
                                                                                                       Query Match
Best Local Similarity 36.8%;
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2 IREDEE--KEGSEQ--
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; ORGANISM: Homo sapiens
US-10-104-047-2343
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 9
US-08-764-870-9
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ORGANISM:
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Sequence 2, Application US/10274990

GENERAL INFORMATION:

APPLICANT: WEI, Ming-Hui et al

TITLE OF INVENTION: ISOLARED HUMAN TRANSPORTER PROTEINS,

TITLE OF INVENTION: NUCLEIC ACID MOLECULES ENCODING HUMAN TRANSPORTER PROTEINS,

TITLE OF INVENTION: NUCLEIC ACID MOLES THEREOF

FILE REFERENCE: CLOOJO18 DIV

CURRENT APPLICATION NUMBER: US/10/274,990

CURRENT FILING DATE: 2000-10-22

FRIOR APPLICATION NUMBER: 09/734,674

PRIOR FILING DATE: 2000-12-13

NUMBER OF SEQ ID NOS: 4

SOFFWARE: FasteSEQ for Windows Version 4.0

SEQ ID NO S.
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APPLICANT: Schroder, Harwaid
APPLICANT: Schroder, Harwaid
APPLICANT: Applacer, Burkhand
APPLICANT: Applacer, Gregor
APPLICANT: Applacer, Gregor
TITLE OF INVENTION: CORYNBEACTERIUM GLUTAMICUM GENES ENCODING NOVEL
TITLE OF INVENTION: PROTEINS
FILE REFERENCE: BGI-129CP
CURRENT APPLICATION NUMBER: US/09/605,703B
CURRENT PILING DATE: 1999-07-08
PRIOR PILING DATE: 1999-07-08
PRIOR PILING DATE: 1999-07-08
PRIOR FILING DATE: 1999-09-03
NUMBER OF SEQ ID NOS: 2934
SEQ ID NO 2760
                                                                                                                                                                                                                                                                                                    Gaps
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Pred. No. 3.6;
6; Mismatches 18; Indels 5.
                                                                                                                                                                                                                                                          Length 1129;
                                                                                                                                                                                                                                                                                                                                                                    1059 EDAEKEEEQSMLAMEDEGTVQLPL----EGHYRDDPSVINISDEMSK 1101
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                                                                                                                                                                                                                                                                                                                                            4 EDEEKEGSEQALSSRDGAPVQHPLALLPSGSYEEDPQTI-LDDHLSR
                                                                                                                                                                                                                                                                                                  6; Mismatches
                                                                                                                                                                                                                                                          Score 72.5;
Pred. No. 3.
TITLE OF INVENTION: AND USES THEREOF
FILE REPERENCE: CL001018
CURRENT APPLICATION NUMBER: US/09/734,674
CURRENT FILING DATE: 2000-12-13
NUMBER OF SEQ ID NOS: 4
SEQ TO NO 2
LENGTH: 1129
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 2760, Application US/09605703B Patent No. 6962989
                                                                                                                                                                                                                                                          20.1%;
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Best Local Similarity 38.3%;
Matches 18; Conservative 6
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APPLICANT: Pompejus, Markus
                                                                                                                                                                                                                                                                            Best Local Similarity 38.3
Matches 18, Conservative
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US-09-734-674-2
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US-10-274-990-2
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                                                                                                                                                                       TYPE: PRT
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FEATURE:
NAME/KEY: DOMAIN
LOCATION: (168)..(441)
OTHER INFORMATION: minimal ligand binding domain
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SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                54 PGCQSPGVGRYSPRSR
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Best Local Similarity, 30.33
Matches 23; Conservative
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US-09-166-265-7
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                                                                                      US-08-980-115-9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 2; Length 441;
    Pacentin Release #1.0, Version #1.30
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30.3%; Pred. No. 3.3;
tive 10; Mismatches
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CURRENT FILING DATE: 1997-11-26
EARLIER APPLICATION NUMBER: 08/74,870
EARLIER PILING DATE: 1996-12-13
EARLIER FILING DATE: 1996-12-13
EARLIER FILING DATE: 1995-12-14
EARLIER FILING DATE: 1995-12-14
EARLIER FILING DATE: 1995-12-13
EARLIER FILING DATE: 1995-12-13
EARLIER FILING DATE: 1995-12-13
NUMBER OF SEQ ID NOS::17
                                                                                                                                                                                                                                                                                                                                       NAME: NAKAMURA, Jackie N
REGISTRATION NUMBER: 35,966
REFERENCE/DOCKET NUMBER: UCAL-246/01US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (650)843-5000
INFORMATION FOR SEQ ID NO: 9:
                                                                                      CLASSIFICATION: 530
RHIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/008,540
RILING DATE: 13-DEC-1995
RICATION NUMBER: US 60/008,543
APPLICATION NUMBER: US 60/008,543
FILING DATE: 13-DEC-1995
RILING DATE: 14-DEC-1995
ATTORNEY/AGENT INFORMATION:
                                              APPLICATION NUMBER: US/08/764,870 FILING DATE: 13-DEC-1996 CLASSIFICATION: 530
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 9, Application US/08980115
Patent No. 626622
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Scanlan, Thomas S. APPLICANT: Baxter, John D. APPLICANT: Fletterick, Robert J. APPLICANT: Wagner, Richard L. APPLICANT: Rushner, Peter J. APPLICANT: Applicant: Applicant: Applicant: West, James W. APPLICANT: West, Brian L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   60 MGCDGASCGSLNMECR 75
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         REFERENCE: UCAL-246/02US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LENGTH: 441 amino acids
TYPE: amino acid
STRANDEDNESS:
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Best Local Similarity 30.3
Matches 23; Conservative
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MOLECULE TYPE: protein
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Sequence 1157, Application US/09949016
; Sequence 1157, Application US/09949016
; Patent No. 6812339
; GENERAL INFORMATION:
APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
TITLE OF INVENTION: WINDMER: 60/241,755
PRIOR PAPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-03
; PRIOR FILING DATE: 2000-10-03
; PRIOR FILING DATE: 2000-09-08
; PRIOR FILING DATE: 2000-09-08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Cummings, Richard T.
APPLICANT: Hermes, Jeffrey D.
APPLICANT: Hornes, Jeffrey D.
APPLICANT: Holler, David E.
APPLICANT: Zhou, Gaochao
TITLE OF INVENTION: ASSAYS FOR NUCLEAR RECEPTOR AGONISTS AND
TITLE OF INVENTION: ANTAGONISTS USING FLUORESCENCE RESONANCE ENERGY TRANSFER
FILE REFERENCE: 2001, CURRENT APPLICANION NUMBER: US/09/166, 265
CURRENT FILING DATE: 1998-10-05
                                                                                                                                                    10 EVREEEKKEEVAEA----EGAPELNGGPQHALPSSSSYTDLSRSSSPPSLLDQ-----LQ 59
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                                                                                                                1 OIREDEEKEGSEQALSSRDGAPVQH--PLALLPSGSYEE-----DPQTILDDHLSRVLKT 53
                                                         17;
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      Length 441;
                                                         Indels
      DB 2;
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Query Match
19.0%; Score 68.5; Di
Best Local Similarity 30.3%; Pred. No. 3.3;
Matches 23; Conservative 10; Mismatches
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 7
LENGTH: 441
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 7, Application US/09166265
Patent No. 6689574
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-----LSRSSSPPSRLD 56

23;

Length 441;

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GENERAL INFORMATION:
APPLICANT Homburger et al.
APPLICANT Homburger et al.
TITLE OF INVENTION: Nucleic acids and proteins of Drosophila melanogaster
TITLE OF INVENTION: Nucleic acids and proteins of Drosophila melanogaster
FILE REFERENCE: File Reference: 7326-094
CURRENT APPLICATION NUMBER: US/09/270,767
KURRENT FILING DATE: 1999-03-17
NUMBER OF SEQ ID NOS: 62517
SOFTWARE: PatentIN Ver. 2.0
SEQ ID NO 40580
LENGTH: 90
                                                                                                                                                                                                   1 QIREDERKEGSEQALSSRDGAPVQH--PLALLPSGSYEEDPQTILDDHLSRVLKTP---
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                                                                                                                                                                                                                                                     10 EVREEEEKEEVAEA----EGAPELNGGPQHALPSSSYTD-
                                         Score 66.5; DB 2;
Pred. No. 5.9;
8; Mismatches 24;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ORGANISM: Drosophila melanogaster
                                                                                                                                                                                                                                                                                                                                                                 55 ----GCQSPGVGRYSPRSR 69
                                                                                                                                                                                                                                                                                                                                                                                                                                        57 QLQMGCDGASCGSLNMECR 75
                                         18.5%;
30.4%;
                                                                                                                        24; Conservative
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                                         Query Match
Best Local Similaritý
Matches 24; Conserval
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Sequence 1000, Application US/09976594

Patent 1000, Application US/09976594

Sequence 1000, Application US/09976594

Seneral Information:
APPLICANT: Buchbinder, Jenny

TITLE OF INVENTION: GENES EXPRESSED IN C3A LIVER CELL CULTURES TREATED WITH STEROIDS FILE REPERENCE: PA-0041 US

CURRENT APPLICATION NUMBER: US/09/976,594

CURRENT PILING DATE: 2000-10-12

PRIOR FILING DATE: 2000-10-12

NUMBER OF SEQ ID NOS: 1143

SOFTWARE: PERL Program

SEQ ID NO 1000

LENGTH: 441
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APPLICANT:
APPLICANT:
ABRUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
TITLE OF INVENTION:
ABRUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
TITLE OF INVENTION:
ABRUGINOSE:
107196.136
CURRENT APPLICATION NUMBER:
CURRENT FILING DATE:
1999-02-18
PRIOR APPLICATION NUMBER:
1999-02-18
PRIOR APPLICATION NUMBER:
1999-02-18
PRIOR APPLICATION NUMBER:
1999-07-27
NUMBER OF SEQ ID NOS:
33142
SEQ ID NO 17434
LENGTH: 595
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; OTHER INFORMATION: Incyte ID No. 6673549 678004CD1
US-09-976-594-1000
                                                                                                                                                                                                                                             DB 2;
                                                                                                                                                                                                                                    19.0%; Score 68.5; DE ilarity 30.3%; Pred. No. 3.9; Conservative 10; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 17434, Application US/09252991A Patent No. 6551795 GENERAL INFORMATION: APPLICANT: Marc J. Rubenfield et al.
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Best Local Similarity
Matches 23; Conserv
SEQ ID NO 11597
LENGTH: 500
TYPE: PRT
ORGANISM: Human
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US-09-252-991A-17434
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Sequence 116, App
Sequence 8, Appli
Sequence 36, Appli
Sequence 35, Appl
Sequence 270, App
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670, App
227845,
                                                         April 20, 2006, 15:57:53 ; Search time 23.971 Seconds (without alignments) 1202.714 Million cell updates/sec
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(cgn2_6/ptodata/1/pubpaa/US07_PUBCOMB.pep:*

cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep:*

cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep:*

cgn2_6/ptodata/1/pubpaa/US10A_PUBCOMB.pep:*

cgn2_6/ptodata/1/pubpaa/US10A_PUBCOMB.pep:*

cgn2_6/ptodata/1/pubpaa/US10B_PUBCOMB.pep:*
GenCore version 5.1.7
Copyright (c) 1993 - 2006 Biocceleration Ltd
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US-10-425-115-205230
US-10-425-114-56486
US-09-920-804-2
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US-11-061-825-2
US-110-450-753-33467
US-10-425-115-244452
US-09-738-626-3956
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-10-425-115-262984
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                                                                                                                                                                          1867569 segs, 417829326 residues
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US-09-734-674-2
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                                        OM protein - protein search, using sw model
                                                                                                                                                      Gapop 10.0 , Gapext 0.5
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Maximum DB seq length: 2000000000
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Match Length DB
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Et Tissue Sarcoma, Compositions
Soft Tissue Sarcoma Modulators
                                                                                                                                      Sequence 33468, A
Sequence 975, App
Sequence 270243,
Sequence 14535, Ap
                                                                                                                                                                     Sequence 8055, Ap
Sequence 14535, A
Sequence 118666,
                                   Appl
                                                                 352367,
                                                                               110790,
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                      Sequence
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Sequence
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Sequence 2
Sequence 1
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Sequence
US-10-104-047-2343

US-10-486-977-11

US-10-690-880-39

US-10-690-880-39

US-10-496-377-7

US-10-496-377-7

US-10-425-115-352367

US-10-425-115-35317

US-10-767-701-38282

US-10-450-763-384496

US-10-450-763-34489

US-10-450-763-3468

US-10-450-763-3468

US-10-369-493-975

US-10-369-493-975

US-10-369-493-975
                                                                                                                                                                                                                                                     RESULT 1
US-10-723-860-1797
Sequence 1797, Application US/10723860
Publication No. US20040253606A1
GENERAL INPORMATION:
APPLICANT: Aziz, Natasha
APPLICANT: Ginsburg, Wendy M.
APPLICANT: Ginsburg, Wendy M.
APPLICANT: Ginsburg, Wendy M.
APPLICANT: Ginsburg, Wendy M.
APPLICANT: Ginsburg, Wendy M.
APPLICANT: Ginsburg, Wendy M.
APPLICANT: Ginsburg, Wendy M.
APPLICANT: Ginsburg, Wendy M.
APPLICANT: Solonik, Albert
TITLE OF INVENTION: Methods for Screening fc
TITLE OF INVENTION: Methods for Screening fc
CURRENT APPLICATION NUMBER: US/10/723,860
CURRENT FILING DATE: 2003-11-26
NUMBER OF SEQ ID NOS: 8393
SOPTWARE: PATENTIN VERSION 3.2
SEQ ID NO 1797
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 19.0
18.9
18.8
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n US/1 3606A1	ndy M ert ods o

; ORGANISM: Homo sapiens US-10-723-860-1797 LENGTH: 843 TYPE: PRT

ö 92.2%; Score 332; DB 5; Length 843; 89.9%; Pred. No. 5.3e-30; Indels 1; Mismatches 3; Indels Ouery Match Dest Local Similarity 89.3.
Best Local Similarity 89.3.

456 1 QIREDEEKEGSEQALSSRDGAPVQHPLALLPSGSYEEDPQTILDDHLSRVLKTPGCQSPG

61 VGRYSPRSR 69

Sequence 116, Application US/10751736
Publication No. US20040265230A1
GENERAL INFORMATION:
APPLICANT: Wyeth
APPLICANT: Wyeth
APPLICANT: West
APPLICANT: Liu, Wei
APPLICANT: Liu, Wei
APPLICANT: Liu, Wei
APPLICANT: Liu, Wei
APPLICANT: Liu, Wei
APPLICANT: Liu, Wei
APPLICANT: Liu, Wei
APPLICANT: Liu, Wei
APPLICANT: Liu, Wei
APPLICANT: Liu, Wei
APPLICANT: CANCERS
FIFTE REFERENCE: AM10929-7 (031896-002000)
CURRENT APPLICATION NUMBER: 2003-01-06
CURRENT FILING DATE: 2003-01-06 RESULT 2 US-10-751-736-116

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APPLICANT: Wyeth
APPLICANT: O'Toole, Margot
APPLICANT: O'Toole, Margot
APPLICANT: Liu, Well
ITILE OF INVENTION: COMPOSITIONS AND METHODS FOR DIAGNOSING AND TREATING AUTOIMMUNE
TITLE OF INVENTION: DISEASES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ų.
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35.8%; Pred. No. 3.1e-05;
tive 10; Mismatches 21;
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Best Local Similarity 35.8%; Pred. No. 3.3e-05;
Matches 34; Conservative 10; Mismatches 21;
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FILE REFERENCE: 031896-023000 (AM101331L)
CURRENT APPLICATION NUMBER: US/10/786,720
CURRENT FILING DATE: 2004-02-26
NUMBER OF SEQ ID NOS: 21135
SOFTWARE: PatentIn version 3.2
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; Publication No. US20040041382A1
; GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Padigaru, Muralidhara
APPLICANT: Spytek, Kimberly A.
APPLICANT: Shenoy, Suresh G.
APPLICANT: Taupler Jr., Raymond
APPLICANT: Pena, Carol E.A.
APPLICANT: Li, Li
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Gusev, Vladimir Y.
Ji, Weizhen
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Gangolli, Esha A.
                                                                                                                                 Query Match 33.9
Best Local Similarity 35.8
Matches 34; Conservative
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; SEQ ID NO 36
; LENGTH: 826
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-786-720-36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: PRT
CORGANISM: Homo sapiens
US-10-786-720-35
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APPLICANT: 0'Toole, Wei
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR DIAGNOSING AND TREATING AUTOIMMUNE
TITLE OF INVENTION: DISEASES
FILE REFERENCE: 031896-023000 (AMI01331L)
CURRENT APPLICATION NUMBER: US/10/786,720
CURRENT FILING DATE: 2004-02-26
NUMBER OF SEO ID NOS: 21135
SOFTWARE: Patentin version 3.2
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Sequence 8, Application US/09798831
Patent No. US20010052137A1
GENERAL INFORMATION:
TITLE OF INVENTION: AXIN DOMAIN-LIKE POLYPEPTIDE INHIBITORS OF GLYCOGEN
TITLE OF INVENTION: SYNTHASE KINASE 3(BETA) ACTIVITY AND ACTIVATORS OF WIT
TITLE OF INVENTION: SIGNALING
FILE REFERENCE: 209596.0391/306U1
CURRENT APPLICATION NUMBER: US/09/798,831
CURRENT FILING DATE: 2001-03-01
PRIOR APPLICATION NUMBER: US 60/186,141.
PRIOR FILING DATE: 2000-03-01
NUMBER OF SEQ ID NOS: 12
SOFTWARE: PatentIn Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            397 QIREDEEREGSELTLNSREGAPTQHPLSLLPSGSYEEDPQTILDDHLSRVLKTPGCQSPG 456
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ---SYEEDPOTILD 44
                                                                                                                                                                                                                                                                                                                   1 QIREDEEKEGSEQALSSRDGAPVQHPLALLPSGSYEEDPQTILDDHLSRVLKTPGCQSPG 60
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                                                                                                                                                                                                                                                                         Gaps
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PRIOR APPLICATION NUMBER: US Provisional Application 60/438,000 PRIOR FILING DATE: 2003-01-06 NUMBER OF SEC ID NOS: 54873 SOFTWARE: Patentin version 3.2 SEC ID NO 116 LENGTH: 843
                                                                                                                                                                                                                          Query Match 92.2%; Score 332; DB 5; Length 843; Best Local Similarity 89.9%; Pred. No. 5.3e-30; Matches 62; Conservative 4; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 43.9%; Score 158; DB 3; Length 842; Best Local Similarity 40.0%; Pred. No. 1.8e-09; Matches 34; Conservative 14; Mismatches 11; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    7 EKEGSEQALSSRDGAPVOHPLALLPSG----
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APPLICANT: Wyeth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ORGANISM: Xenopus laevis
                                                                                                                                                                                                                                                                                                                                                                                                                                                       457 VGRYSPRSR 465
                                                                                                                                                                                                                                                                                                                                                                                                               61 VGRYSPRSR 69
                                                                                                                                                             ORGANISM: Homo sapiens
                                                                                                                                                                                  US-10-751-736-116
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 31.5%; Score 113.5; DB 4; Length S
Best Local Similarity 34.4%; Pred. No. 0.00035;
Matches 33; Conservative 10; Mismatches 22; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    33 GSYEEDPOTILDDHLSRVLKTPGCQSPGVGRYSPRS
PRIOR APPLICATION NUMBER: US 09/543,771
PRIOR FILING DATE: 2000-04-05
PRIOR APPLICATION NUMBER: US 09/229,319
PRIOR PILING DATE: 1999-01-13
PRIOR PLING-DATE: 1999-01-13
PRIOR PELICATION NUMBER: US 60/071,449
PRIOR APPLICATION NUMBER: US 60/105,511
PRIOR PILING DATE: 1998-10-23
NUMBER OF SEQ ID NOS: 109
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Allen, Kristina M.
APPLICANT: Anisowicz, Anthony
APPLICANT: Bhat, Bheem
APPLICANT: Damagnez, Veronique
APPLICANT: Robinson, John
APPLICANT: Robinson, John
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Best Local Similarity 34.44
Matches 33; Conservative
                                                                                                                                                                                                                                                                                                                                                                                     TYPE: PRT ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: PRT
ORGANISM: Homo sapiens
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                                                                                                                                                                                                                                                                                                                 SEQ ID NO 91
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TITLE OF INVENTION: No. US20040043382Alel Proteins and Nucleic Acids Encoding Same
FILE REFERENCE: 21402-290C
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Publication No. US20030219793A1
GENERAL INFORMATION:
APPLICANT: JOHN F. CATULLI et al.
TITLE OF INVENTION: THE HIGH BONE MASS GENE OF 11q13.3
FILE REFERENCE: 032796-021
CURRENT APPLICATION NUMBER: US/10/374,979
CURRENT FILING DATE: 2003-03-04
PRIOR PILING DATE: 2000-04-05
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35.4%; Pred. No. 4.1e-05;
ative 10; Mismatches 21
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PRIOR FILING DATE: 2001-04-13
PRIOR FILING DATE: 2001-04-13
PRIOR FILING DATE: 2001-04-13
PRIOR FILING DATE: 2001-03-08
PRIOR PLING DATE: 2001-03-08
PRIOR APPLICATION NUMBER: USSN 60/274,191
PRIOR APPLICATION NUMBER: USSN 60/294,354
PRIOR PLING DATE: 2001-07-10
PRIOR PRIOR DATE: 2001-03-30
PRIOR PLING DATE: 2001-03-30
PRIOR PLING DATE: 2001-03-30
PRIOR PLING DATE: 2001-03-30
PRIOR PLING DATE: 2001-03-30
PRIOR PLING DATE: 2001-03-30
PRIOR PLING DATE: 2001-03-30
PRIOR PLING DATE: 2001-03-30
PRIOR PLING DATE: 2001-05-31
PRIOR PLING DATE: 2001-05-31
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CURRENT FILING DATE: 2002-03-07
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PRIOR APPLICATION NUMBER: USSN 60/274,322
PRIOR FILING DATE: 2001-03-08
                                       Chernev,, Velizar T. Pernandes, Elma R.
                                                                                                        Casman, Stacie J.
Malyankar, Uriel M.
Gerlach, Valerie
                                                                                                                                                                                                                                                                                    Steven K
          Xiaojia Sasha
                                                                                                                                                                                                                                                                                                                                                                                                                         Alsobrook, John P.
Lepley, Denise M.
Rieger, Daniel K.
                                                                                                                                                                                                                                                Anderson, David W.
                                                                                                                                                                                                                                                                                                                 Catterton, Elina
Leite, Mario W.
Zhong, Haihong
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Best Local Similarity 35.4
Matches 34; Conservative
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SEQ ID NO 270
LENGTH: 912
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US-10-092-900A-270
                                                                                                                                                                                                                                                                                    Spaderna,
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Best Local Similarity 34.4%; Pred. No. 0.00035; Matches 33; Conservative 10; Mismatches 22; Indels
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                                                                    ---EGSEQALSSRDGAPVQHPLALLPS
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                                                                                                                                                             33 GSYEEDPOTILDDHLSRVLKTPGCQSPGVGRYSPRS
                                                                                                                                                                                                                                                                                                                  Sequence 670, Application US/10477173
Publication No. US20050070699A1
GENERAL INFORMATION:
APPLICANT: Genome Therapeutics Corporation and APPLICANT: Yaworsky, Paul
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CURRENT APPLICATION NUMBER: US/10/425,115
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 369326
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQ ID NOS: 1086
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR PILING DATE: 2001-05-17
PRIOR APPLICATION NUMBER: US 60/353,058
PRIOR FILING DATE: 2002-02-01
PRIOR PLICATION NUMBER: US 60/361,293
PRIOR FILING DATE: 2002-03-04
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT FILING DATE: 2003-11-10
PRIOR APPLICATION NUMBER: US 60/290,071
PRIOR FILING DATE: 2001-05-11
PRIOR APPLICATION NUMBER: US 60/291,311
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Morales, Arturo J.
Graham, James R.
Anisowicz, Anthony
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ORGANISM: Homo sapiens
                                                                    3 REDEEK-----
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Best Local Similarity
Matches 33; Conserv
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                                                                                       APPLICANT: Yaworsky, Paul
APPLICANT: Bex, Frederick J. III
APPLICANT: Box, Frederick J. III
APPLICANT: Bodine, Peter Van Nest
TITLE OF INVENTION: Transgenic Animal Model of Bone Mass Modulation
FILE REFERENCE: 032796-212
CURRENT APPLICATION NUMBER: US/10/477,238A
CURRENT FILING DATE: 2003-11-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Yaworsky, Paul
APPLICANT: Bex, Frederick J. III
APPLICANT: Bodine, Peter Van Nest
TITLE OF INVENTION: Transgenic Animal Model of Bone Mass Modulation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 900;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 5;
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CURRENT APPLICATION NUMBER: US/10/680,287A
CURRENT FILING DATE: 2003-10-08
PRIOR FILING DATE: 2001-05-13
PRIOR FILING DATE: 2001-05-11
PRIOR FILING DATE: 2001-05-11
PRIOR FILING DATE: 2001-05-17
PRIOR FILING DATE: 2001-05-17
PRIOR FILING DATE: 2001-05-17
PRIOR FILING DATE: 2001-05-17
PRIOR FILING DATE: 2002-02-01
PRIOR FILING DATE: 2002-03-04
PRIOR FILING DATE: 2002-03-04
NUMBER OF SEQ ID NOS: 812
NUMBER OF SEQ ID NOS: 812
SEQ ID NO 670
LENGTH: 900
                                                                                                                                                                                                                                                     CURRENT FILING DATE: 2003-11-10
PRIOR APPLICATION NUMBER: US 60/290,071
PRIOR PILING DATE: 2001-05-11
PRIOR APPLICATION NUMBER: US 60/291,311
PRIOR APPLICATION NUMBER: US 60/353,058
PRIOR FILING DATE: 2002-02-01
PRIOR PLING DATE: 2002-02-01
PRIOR PLING DATE: 2002-03-04
PRIOR PILING DATE: 2002-03-04
NUMBER OF SEQ ID NOS: 812
SOFTWARE: FastSEQ for Windows Version 4.0
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US-10-477-238A-670
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US-10-680-287A-670
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LENGTH: 900
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Sequence 227845, Application US/10425115
Publication No. US20040214272A1
GENERAL INFORMATION:
APPLICANT: La Rosa, Thomas J.
APPLICANT: Zhou, Yihua
APPLICANT: Zhou, Yihua
APPLICANT: Character Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control of Control
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Gaps

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Sequence 205230, Application US/10425115
Publication No. US20040214272A1
GAPPLICANT: INFORMATION:
APPLICANT: Kovalic, David K.
APPLICANT: Cao, Yongwei
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants
FILE OF INVENTION: Plants
CURRENT APPLICATION NUMBER: US/10/425,115
CURRENT APPLICATION NUMBER: US/10/425,115
CURRENT PILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 369326
SECOND NO 205230
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With TITLE OF INVENTION: Plants PILE REFERENCE: 38-21 (33.22) B CURRENT APPLICATION NUMBER: US/10/425,115 CURRENT APPLICATION NUMBER: US/203-04-28 NUMBER OF SEQ ID NOS: 369326 SEQ ID NOS: 369326 LENGTH: 175
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                                                                                                                                                                                                                                                                                                                                                Score 73; DB 4; Length 175; Pred. No. 2.7; 8; Mismatches 28; Indels
                                                                                                                                                                                                                                                                 , OTHER INFORMATION: Clone ID: MRT4577_139392C.1.pep
US-10-425-115-227847
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, OTHER INFORMATION: Clone ID: MRT4577_118759C.1.pep
US-10-425-115-205230
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                                                                                                                                                                                                                         ORGANISM: Zea mays
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ORGANISM: Zea mays
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APPLICANT: La Rosa, Thomas J. APPLICANT: Ea Royalice, David K. APPLICANT: Shou, Yihua APPLICANT: Zhou, Yihua APPLICANT: Cao, Yongwei APPLICANT: Cao, Yongwei APPLICANT: Barbazuk, Brad APPLICANT: Barbazuk, Brad APPLICANT: Barbazuk, Brad APPLICANT: Li, Ping APPLICANT: Li, Ping III SO INVENTION: Rice Nucleic Acid Molecules and Other Molecules Associated With TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement FILE REFERENCE: 38-21(5321)8

CURRENT APPLICATION NUMBER: 20/10/437,963

CURRENT PILING DATE: 2003-05-14
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                                                                                                                                                                                                                                                                         Query Match 20.8%; Score 75; DB 4; Length 218; Best Local Similarity 29.5%; Pred. No. 2.1; Matches 23; Conservative 6; Mismatches 29; Indels
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US-10-437-963-124794
                                                                                                                                                                                                     ; OTHER INFORMATION: Clone ID: MRT4577_139390C.1.pep
US-10-425-115-227845
                                                                                                                                LOCATION: (1)..(218)
OTHER INFORMATION: unsure at all Xaa locations
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Publication No. US20040214272A1
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       145 RLORVCSSPAPFXAGCSS 162
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APPLICANT: Kovalic, David K.
APPLICANT: Zhou, Yihua
APPLICANT: Cao, Yongwei
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Matches 23; Conservative
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                 LENGTH: 218
TYPE: PRT
ORGANISM: Zea mays
                                                                                                     NAME/KEY: unsure
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LENGTH: 1487
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Sequence 20102, A Sequence 20101, A Sequence 6198, Ap Sequence 901, App Sequence 5344, Ap Sequence 6244, Ap Sequence 6268, Ap
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Sequence 6507, Ap
Sequence 2474, Ap
Sequence 11072, A
Sequence 11072, A
Sequence 11034, Ap
Sequence 17930, A
Sequence 1960, Ap
Sequence 60, Appl
Sequence 61937, A
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Sequence 2343, Ap
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Sequence 20102, A
                                                                                       April 20, 2006, 16:00:23 ; Search time 3.82322 Seconds (without alignments) 794.148 Million cell updates/sec
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1: /SIDS5/ptodata/2/pubpaa/US08 NEW PUB.pep:*

2: /SIDS5/ptodata/2/pubpaa/US07 NEW PUB.pep:*

3: /SIDS5/ptodata/2/pubpaa/US07 NEW PUB.pep:*

4: /SIDS5/ptodata/2/pubpaa/PCT NEW PUB.pep:*

5: /SIDS5/ptodata/2/pubpaa/US10 NEW PUB.pep:*

7: /SIDS5/ptodata/2/pubpaa/US10 NEW PUB.pep:*

8: /SIDS5/ptodata/2/pubpaa/US10 NEW PUB.pep:*

8: /SIDS5/ptodata/2/pubpaa/US10 NEW PUB.pep:*
GenCore version 5.1.7
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US-11-096-568A-20102
US-11-096-568A-20101
US-11-188-298-6198
US-11-188-298-11473
US-11-188-298-11473
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-11-188-298-17930
-11-188-298-1960
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US-10-204-639-64
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US-11-072-512-2474
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                                                                                                                                                                                                                                                                                            Total number of hits satisfying chosen parameters:
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                                                                                                                                                                                                                                                                  225428 segs, 44002918 residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SUMMARIES
                                                                                                                                                                                                                                                                                                                                                                          Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                            OM protein - protein search, using sw model
                                                                                                                                                                                                                                   Gapop 10.0 , Gapext 0.5
                                                                                                                                                                                                                                                                                                                            Minimum DB seq length: 0
Maximum DB seq length: 200000000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query
Match Length DB
                                                                                                                                                   US-09-587-574-4
360
                                                                                                                                                                                                                    BLOSUM62
                                                                                                                                                      Title:
Perfect score:
                                                                                                                                                                                                                    Scoring table:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        113.5
69.5
65
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   63.5
63.5
63.5
62.5
62.5
61.5
61.5
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                                                                                                                                                                                       Sequence:
                                                                                                                                                                                                                                                                  Searched:
                                                                                            Run on:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Result
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Sequence Sequence Sequence

US-11-096-568A-11937 US-11-096-568A-12944 US-11-096-568A-12943 US-10-467-657-2944

58 16.1 297 7 US-11-096-568A-12942 Sequence 12942, A 58 16.1 418 6 US-10-312-954-4 Sequence 13444, A 57.5 16.0 200 7 US-11-188-298-1344 Sequence 13444, A 57.5 16.0 202 7 US-11-188-298-13134 Sequence 19134, A 57.5 16.0 202 7 US-11-188-298-20151 Sequence 19134, A 57.5 16.0 202 7 US-11-188-298-21517 Sequence 19134, A 57.5 16.0 202 7 US-11-188-298-21517 Sequence 20480, A 57.5 16.0 204 7 US-11-188-298-2050 Sequence 20480, A 57.5 16.0 204 7 US-11-188-298-20490 Sequence 20480, A 57.5 16.0 5435 6 US-10-453-372-1002 Sequence 192, Apple 57.5 15.8 513 6 US-10-31-826A-192 Sequence 192, Apple 57.5 15.8 513 6 US-10-31-818-192 Sequence 192, Apple 57.5 15.8 513 6 US-10-31-156 Sequence 192, Apple 57.5 15.8 513 6 US-10-995-561-566 Sequence 192, Apple 57.5 15.8 513 6 US-10-995-561-566 Sequence 192, Apple 57.5 15.8 513 6 US-10-995-561-566 Sequence 192, Apple 57.5 15.8 513 6 US-10-995-561-566 Sequence 192, Apple 57.5 15.8 513 6 US-10-995-561-566 Sequence 192, Apple 57.5 15.8 513 6 US-10-995-561-566 Sequence 192, Apple 57.5 15.8 18.7 US-11-120-955-3 Sequence 192, Apple 57.5 15.8 18.7 US-11-188-298-18071 Sequence 24982, Apple 56.5 15.7 15.8 7 US-11-188-298-18071 Sequence 24982, Apple 56.5 15.7 15.8 7 US-11-188-298-18071 Sequence 24982, Apple 56.5 15.7 218 7 US-11-1096-568A-24982	SULT 1  -10-501-035-215  Sequence 215, Application US/10501035  Sequence 215, Application US/10501035  Sequence 215, Application US/10501035  Sequence 215, Application US/10501035  Sequence 215, Application US/10501035  SEQUENCE 215  GENERAL INFORMATION: DEBATIFICATION OF POLYMUCLEOTIDES AND POLYPEPTIDE FOR PREDICTIN TITLE OF INVENTION: ACTIVITY OF COMPOUNDS THAT INTERACT WITH PROTEIN TYROSINE KINATITLE OF INVENTION: ADD/OR PROTEIN TYROSINE KINASE PATHWAYS  FILE REFERENCE: DO1045 FOT CURRENT APPLICATION NUMBER: US/10/501,035  CURRENT APPLICATION NUMBER: US 60/350,061  PRIOR APPLICATION NUMBER: US 60/350,061  RIOR APPLICATION NUMBER: US 60/350,061  RIOR APPLICATION NUMBER: US 60/350,061  RIOR APPLICATION NUMBER: US 60/350,061  NUMBER OF SEQ ID NOS: 795  SEQ ID NO 215  LENGTH: 900  TYPE: PRT  ORGANISM: Homo sapiens  -10-501-035-215	Query Match  31.5%; Score 113.5; DB 6; Length 900;  Best Local Similarity 34.4%; Pred. No. 1.4e-05;  Matches 33; Conservative 10; Mismatches 22; Indels 31; Gaps 3;  3 REDEEK
266 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5	RESULT 1 US-10-501-035-2 Sequence 215, Sequence 215, Publication Ni GENERAL INFORM APPLICANT: B. TITLE OF INV TITLE OF INV TITLE OF INV TITLE OF INV TITLE OF INV TITLE OF INV TITLE OF INV TITLE OF INV TITLE OF INV TITLE OF INV TITLE OF INV TITLE OF INV TITLE OF INV TITLE OF INV TITLE OF INV SECUENCE APPLIC RAID NO 215 SEQ ID NO 215 LENGTH: 900 TIPLE PRI SEQ ID NO 215 LENGTH: 900 TIPLE ORGANISM: H US-10-501-035-2.	Ouery Match Best Local Simil Batches 33; C Oy 3 REDE Db 440 REAE Oy 33 GSYE OS 11-072-512-2343; A SEQUENCE 2343; ``
Sequence 20101. Application US/11096568A

Publication No. US20060048240A1

GENERAL INFORMATION:

APPLICANI: Alexandrow, Nickolai et al.

TITLE OF INVENTION: Sequence-Determined DNA Fragments and Corresponding Polypeptide

TITLE OF INVENTION: Therby

TITLE OF INVENTION: Therby

TITLE OF INVENTION: 1905-1592PUSC

CURRENT APPLICATION NUMBER: US/11/096,568A

CURRENT FILING DATE: 2005-04-01

NUMBER OF SEQ ID NOS: 34471

SEQ ID NO 20101
                                      APPLICANT: Alexandrov, Nickolai et al.
TITLE OF INVENTION: Sequence-Determined DNA Fragments and Corresponding Polypeptide
TITLE OF INVENTION: Therby
FILE REFERENCE: 2750-1592PUS2
CURRENT APPLICATION NUMBER: US/11/096,568A
CURRENT FILING DATE: 2005-04-01
NUMBER OF SEQ ID NOS: 34471
  ب
.
   Gaps
   14;
  14;
  Length '293;
  Query Match
18.1%; Score 65; DB 7; Length 300;
Best Local Similarity 44.2%; Pred. No. 2.9;
Matches 19; Conservative 2; Mismatches 8; Indels
   8; Indels
   48 PGRHP----PSSYQQNGSYESDDRECTPHRIHDDILDSILKTP 86
  22 PVQHPLALLPS----GSYEED-----PQTILDDHLSRVLKTP 54
   APPLICANT: Abad, Mark S. et al.
TITLE ON INVENTION: GENES AND USES FOR PLANT IMPROVEMENT
FILE REFERENCE: 38-21(53452)B
CURRENT APPLICATION NUMBER: US/11/188,298
CURRENT PILING DATE: 2005-07-22
PRIOR REPLICATION WUMBER: 60/592,978
PRIOR FILING DATE: 2004-07-31
NUMBER OF SEQ ID NOS: 22569
   22 PVQHPLALLPS----GSYEED-----PQTILDDHLSRVLKTP
  Query Match 18.1%; Score 65; DB 7;
Best Local Similarity 44.2%; Pred. No. 2.8;
Matches 19; Conservative 2; Mismatches
   LOCATION: (1). (293)
CTHER INFORMATION: Ceres Seq. ID no. 12376899
US-11-096-568A-20102
   ; LOCATION: (1). 7 (300)
; OTHER INFORMATION: Ceres Seq. ID no. 12376898
US-11-096-568A-20101
   ; Sequence 6198, Application US/11188298
; Publication No. US20060075522A1
; GENERAL INFORMATION:
   TYPE: PRT ORGANISM: Zea mays subsp. mays
   ORGANISM: Zea mays subsp. mays
US20060048240A1
  NAME/KEY: misc feature
   NAME/KEY: misc feature
   RESULT 5
US-11-096-568A-20101
   RESULT 6
US-11-188-298-6198
  SEQ 1D NO 20102
   SEQ ID NO 6198
   TYPE: PRT
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   q
   APPLICANT: Alexandrov, Nickolai et al.
APPLICANT: Alexandrov, Nickolai et al.
TITLE OF INVENTION: Sequence-Determined DNA Fragments and Corresponding Polypeptides
TITLE OF INVENTION: Sequence-Determined Corresponding Polypeptides
TITLE OF INVENTION: Therby
FILE REFERENCE: 2750-1592PUS2
CURRENT APPLICATION NUMBER: US/11/096,568A
CURRENT FILING DATE: 2005-04-01
NUMBER OF SEQ ID NOS: 34471
  2 IREDEE--KEGSEQ-----ALSS-----RDGAPVQHPLALLPSGSYEEDPQTI 42
  Gaps
   Query Match
19.3%; Score o.v., ..., Best Local Similarity 28.8%; Pred. No. 4;
Matches 23; Conservative 15; Mismatches 11; Indels 31;
  14;
   Length 257;
  12 PGRHP----PSSYQQNGSYESDDRECTPHRIHDDILDSILKTP 50
   22 PVQHPLALLPS----GSYEED-----PQTILDDHLSRVLKTP 54
  Query Match

18.1%; Score 65; DB 7;
Best Local Similarity 44.2%; Pred. No. 2.4;
Matches 19; Conservative 2; Mismatches
  NAME/KEY: misc_feature
i LOCATION: (1)...(257)
cyter INFORMATION: Ceres Seq. ID no. 12376900
US-11-096-568A-20103
   APPLICANT: APPLICANT: APPLICANT: APPLICANT: MAGAHARI, KENJI
APPLICANT: MASUHO, YASUHIKO
TITLE OF INVENTION: Novel full length cDNA
FILE REFERENCE: 084335-0191
CURRENT APPLICATION NUMBER: US 60/350,978
PRIOR APPLICATION NUMBER: US 60/350,978
PRIOR FILING DATE: 2005-01-25
PRIOR FILING DATE: 2001-12
PRIOR FILING DATE: 2001-11-05
NUMBER OF SEQ ID NOS: 4096
SOFTWARE: PAEGNIT VEY: 2.1
SEQ ID NO 2343
LENGTH: 1034
  Sequence 20103, Application US/11096568A Publication No. US20060048240A1
   ; Sequence 20102, Application US/11096568A
   1::|| ::|| || || || 702 LNNHLGKATEPLRSPSCSSP 721
   43 LDDHLSRV---LKTPGCQSP 59
                  OTSUKA, KAORU
NAGAI, KEIZCHI
IRIE, RYOTARO
TAMECHIKA, ICHIRO
SEKI, NAOHIKO
YOSHIKAWA, TSUTOWU
OTSUKA, MOTOYUKI
  ORGANISM: Zea mays subsp.
  TYPE: PRT
ORGANISM: Homo sapiens
US-11-072-512-2343
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US-11-096-568A-20103
  US-11-096-568A-20102
   SEQ ID NO 20103
LENGTH: 257
  APPLICANT:
APPLICANT:
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APPLICANT:
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Gaps

3

12, Indels

Length 352;

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APPLICANT: Abad, Mark S. et al.
TITLE OF INVENTION: GENES AND USES FOR PLANT IMPROVEMENT
TITLE OF INVENTION: GENES AND USES FOR PLANT IMPROVEMENT
TITLE REPRENCE: 38-21(5345)8
CURRENT APPLICATION NUMBER: US/11/188,298
CURRENT FILING DATE: 2005-07-22
PRIOR PEDLICATION NUMBER: 60/592,978
PRIOR FILING DATE: 2004-07-31
NUMBER OF SEQ ID NOS: 22569
                     APPLICANT: Abad, Mark S. et al.
TITLE OF INVENTION. GENES AND USES FOR PLANT IMPROVEMENT
FILE REFERENCE: 38-21(53452)8
CURRENT APPLICATION NUMBER: US/11/188,298
CURRENT FILING DATE: 2005-07-22
PRIOR APPLICATION NUMBER: 60/592,978
PRIOR FILING DATE: 2004-07-31
SEQ ID NOS: 22569
LENGTH: 352
  APPLICANT: Abad, Mark S. et al.
TITLE OF INVENTION: GENES AND USES FOR PLANT IMPROVEMENT
FILE REFERENCE: 38-21(5345)B
CURRENT APPLICATION NUMBER: US/11/188,298
CURRENT FILING DATE: 2005-07-22
  1 QIREDEEKEGSEQALSSR---DGAPVQHPLALLPSGSYEE 37
  1 OIREDEEKEGSEQALSSR---DGAPVQHPLALLPSGSYEE 37
   k; Score 62.5; DB 7;
k; Pred. No. 7.3;
12; Mismatches 12;
   ; ORGANISM: Oryza sativa (japonica cultivar-group)
US-11-188-298-5344
  Score 62.5; D
Pred. No. 7.3;
  12; Mismatches
  Sequence 6924, Application US/11188298
, Publication No. US20060075522A1
, GENERAL INFORMATION:
  ; Sequence 2268, Application US/11188298
; Publication No. US20060075522A1
; GENERAL INFORMATION:
   PRIOR APPLICATION NUMBER: 60/592,978
PRIOR FILING DATE: 2004-07-31
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   Query Match
Best Local Similarity 32.5%;
Matches 13; Conservative 1:
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Best Local Similarity 32.55
Matches 13, Conservative
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US-11-188-298-6924
  Glycine max
   US-11-188-298-6924
  US-11-188-298-2268
  US-11-188-298-2268
   SEQ ID NO 6924
LENGTH: 352
   SEQ ID NO 2268
   161
   TYPE: PRT
  ORGANISM:
  ò
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   5;
  11 SEQALSSRDGAPVQHPLALLPSGSYEEDPQTILDDHLSRVLKTPGCQSPGVG--RYSPRS 68
   10 GSEQALSSRDGAPVQHPLALLPSGSYEEDPQTILDDHLSRVLKTPGCQSPGVGRY 64
  21 APVQHPLAL---LPSGS----YEBDPQTILDDHLSRVLKTPGCQSPGVGRYS 65
   Sequence 901, Application US/10330773

Publication No. US20060040262A1

GENERAL INFORMATION:
APPLICANT: David W. Morris
APPLICANT: Marc Malandro
FILER REFERENCE: 529452001300
CURRENT APPLICATION NUMBER: US/10/330,773

CURRENT FILING DATE: 2002-12-27

NUMBER OF SEQ ID NOS: 981

SOFTWARE: FastSEQ for Windows Version 4.0

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  APPLICANT: Abad, Mark S. et al.
TITLE OF INVENTION: GENES AND USES FOR PLANT IMPROVEMENT
FILE REFERENCE: 38-21(53422)
CURRENT APPLICATION NUMBER: US/11/188,298
CURRENT FILING DATE: 2005-07-22
PRIOR APPLICATION NUMBER: 60/592,978
PRIOR FILING DATE: 2004-07-31
   17.6%; Score 63.5; DB 7;
32.7%; Pred. No. 3.3;
iive 10; Mismatches 18;
   Score 63.5; DE
Pred. No. 2.8;
   9; Mismatches
   8; Mismatches
  US-11-188-298-11473
; Sequence 11477, Application US/11188298
; Sublication No. US20060075522A1
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; TYPE: PRT
; ORGANISM: Triticum aestivum
US-11-188-298-6198
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   NUMBER OF SEQ ID NOS: 22569
SEQ ID NO 11473
  Best Local Similarity 32.7
Matches 17; Conservative
   Best Local Similarity 35.0
Matches 21, Conservative
   TYPE: PRT
ORGANISM: Homo sapiens
   US-11-188-298-11473
   RESULT 9
US-11-188-298-5344
   US-10-330-773-901
   US-10-330-773-901
   Query Match
  Query Match
  Query Match
  Best Loca
Matches
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Indels

12;

7;

DB

17.1%; Score 61.5;

Query Match

Sequence 5344, Application US/11188298

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Length 352;

DB 7;

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1 QIREDEEKEGSE----QALSSRDGAPVQHPLA-LLPSGSYEEDPQTILDDHLSRVLKTPGC 56
  Sequence 2. Application US/11040488
; Bublication No. US20050271651A1
; GENERAL INFORMATION:
; APPLICATY: WEBB. CAROL
; TITLE OF INVENTION: INHIBITION OF BRIGHT FUNCTION AS A TREATMENT FOR
; TITLE OF INVENTION: EXCESSIVE IMMUNOGLOBULIN PRODUCTION
; TITLE OF INVENTION: EXCESSIVE IMMUNOGLOBULIN PRODUCTION
; TITLE OF INVENTION: EXCESSIVE IMMUNOGLOBULIN PRODUCTION
; TITLE OF INVENTION: EXCESSIVE IMMUNDER: US/11/040,488
; CURRENT FILING DATE: 2005-01-21
; PRIOR APPLICATION NUMBER: 60/538,866
; NUMBER OF SEQ ID NOS: 26
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; SEQ ID NO 2
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   Query Match 16.9%; Score 61; DB Best Local Similarity 33.3%; Pred. No. 36; Matches 23; Conservative 13; Mismatches
   CANT: MASUHO, YASUHIKO
3 OF INVENTION: Novel full length cDNA
REFERENCE: 084335-0191
SNT APPLICATION NUMBER: US/11/072,512
  PRIOR APPLICATION NUMBER: US 60/350,978
PRIOR FILING DATE: 2002-01-25
PRIOR PELICATION NUMBER: JP 2001-379298
PRIOR FILING DATE: 2001-11-05
NUMBER OF SEQ ID NOS: 4096
   Sequence 2474, Application US/11072512
Publication No. US20060029945A1
GENERAL INFORMATION:
   YOSHIKAWA, TSUTOMU
OTSUKA, MOTOYUKI
   YAMAMOTO, JUN-ICHI
ISONO, YUUKO
   KENG
   SHIZUKO
   SATO, HIROYUKI
  WAKAMATSU, AI
   OTSUKA, KAORU
   RYOTARO
  404 MSKGKGKYS 412
   TYPE: PRT
CORGANISM: Homo sapiens
US-11-072-512-2474
   57 OSPGVGRYS 65
  AMP.
ISONO, 1.
YURI
   NAGAHARI,
  SEQ 1D NO 2474
  SOFTWARE
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  APPLICANT: Gary L. Breton
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO BACTEROIDES FRA
TITLE OF INVENTION: NUCLEIC ACID AND THERAPEUTICS
FILE REFERENCE: PATHOD-03DIV2
CURRENT APPLICATION NUMBER: US/11/079,463
CURRENT FILING DATE: 2005-03-14
PRIOR PAPLICATION NUMBER: US 60/128,705
PRIOR PAPLICATION NUMBER: US 09/540,209
PRIOR FILING DATE: 1999-04-09
PRIOR FILING DATE: 2000-04-04
NUMBER OF SEQ ID NOS: 10444
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   6 EEKEGSEQALSSRDGAPVQHPL----ALLPSGSYEEDPQTILDDHLSRVLKTPGCQSPG 60
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   21 APVQHPLAL---LPSGS----YEEDPQTILDDHLSRVLKTPGCQSPGVGRYS 65
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   Sequence 64, A pplication US/10204639
Publication No. US20060063152A1
GENERAL INFORMATION:
GENERAL INFORMATION:
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Osamu Ohara
APPLICANT: Takahiro Nagase
APPLICANT: Daisuke Nakajima
ITILE OF INVENTION: NOVEL GENE AND PROTEIN ENCODED BY THE GENE
FILE REFERENCE: PH-1416-PCT
CURRENT APPLICATION NUMBER: US/10/204,639
CURRENT APPLICATION NUMBER: JP 2000-08-22
PRIOR FILING DATE: 2000-12-22
  17.1%; Score 61.5; DB 6; Length 958; 37.3%; Pred. No. 34;
  Query Match 16.9%; Score 61; DB 7; Length 681; Best Local Similarity 34.5%; Pred. No. 26; Matches 19; Conservative 7; Mismatches 27; Indels
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   27;
    Pred. No. 3.7;
9; Mismatches
   8; Mismatches
  PRIOR APPLICATION NUMBER: JP 2001-095524
PRIOR FILING DATE: 2001-03-29
PRIOR APPLICATION NUMBER: JP 2001-127066
WHENGR FILING DATE: 2001-04-25
NUMBER OF SEQ ID NOS: 140
SOFTWARE: Patentin Ver. 2.1
  Sequence 6507, Application US/11079463 Publication No. US20060073161A1 GENERAL INFORMATION:
       32.7%;
    Best Local Similarity 32.79
Matches 17; Conservative
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Best Local Similarity 37.3
Matches 25; Conservative
  ; LENGTH: 958
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-204-639-64
  850 NGRLHPR 856
  61 VGRYSPR 67
   ORGANISM: B.fragilis
US-11-079-463-6507
   US-11-079-463-6507
  SEQ ID NO 64
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Search completed: April 20, 2006, 16:07:53 Job time : 3.82322 secs .

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April 20, 2006, 15:30:33 ; Search time 4.26121 Seconds (without alignments) 989.497 Million cell updates/sec.
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GenCore version 5.1.7
Copyright (c) 1993 - 2006 Biocceleration Ltd.
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Perfect score:
  Sequence:
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  Run on:
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Listing first 45 summaries

Post-processing: Minimum Match 0% Maximum Match 100%

Minimum DB seq length: 0 Maximum DB seq length: 200000000

Issued\_Patents\_AA:\*

Database :

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

3189, A 3, Appl , Appl , Appl Appl Appl Appl Sequence 1, Appli Sequence 2, Appl Sequence 1, Appli Sequence 22, Appl Sequence 12, Appl Sequence 10957, A Sequence 6495, Ap Sequence 41740, A Sequence 14579, A Sequence 5, Appli Sequence 5, Appli Sequence 5, Appli Sequence 42, Appl Sequence 46, Appl Sequence 48144, A Sequence 10, Appl App] Description Sequence Sequence Sequence Sequence Sequence 4 Sequence 4 Sequence 4 Sequence Sequence Sequence 7 Sequence US-08-890-865A-1 US-08-890-865A-1 US-08-890-865A-1 US-10-464-939-12 US-08-990-865A-21 US-09-949-016-10957 US-09-949-016-10957 US-09-949-016-10957 US-09-949-016-10957 US-09-949-016-10957 US-09-949-016-10957 US-09-990-018-12 US-09-903-018-12 US-09-903-018-12 US-09-903-018-12 US-09-903-018-12 US-09-893-820-42 US-09-893-820-42 US-09-270-767-48144 US-09-270-767-48144 US-09-270-767-48144 SUMMARIES Query Match Length DB Score 56.5 55.5 54.5 54.5 54.5 109.5 102 102 95.5 Result o S

| Sequence 4, Appli<br>Sequence 24, Appli<br>Sequence 10, Appl<br>Sequence 10, Appl<br>Sequence 24, Appl<br>Sequence 24, Appl<br>Sequence 10, Appl<br>Sequence 6574, Appl<br>Sequence 6574, Appl<br>Sequence 2, Appli<br>Sequence 2, Appli<br>Sequence 3, Appli<br>Sequence 8, Appli<br>Sequence 8, Appli<br>Sequence 8, Appli<br>Sequence 8, Appli<br>Sequence 8, Appli<br>Sequence 8, Appli<br>Sequence 3, Appli<br>Sequence 3, Appli<br>Sequence 3, Appli<br>Sequence 3, Appli<br>Sequence 3, Appli                               |            | h 855; els 0; Gaps 0; MYEGRIL 51 ::  :  :  IPEEKII 848                                                                                                                                                                                                           |
|------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|------------|------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|
| 28 54.5 19.7 566 1 US-09-086-662-4 30 53 19.2 309 1 US-09-625-972-24 31 53 19.2 309 1 US-08-655-167A-24 33 53 19.2 309 2 US-08-93-118-10 34 53 19.2 309 2 US-08-93-118-10 35 53 19.2 309 2 US-08-627-820-24 36 53 19.2 309 2 US-09-468-432-10 36 53 19.2 309 2 US-09-468-432-10 37 53 19.2 309 2 US-09-468-432-10 38 53 19.2 309 2 US-09-468-432-10 39 53 19.2 359 2 US-09-949-016-6574 40 52.5 19.0 1332 2 US-09-949-12 41 52.5 19.0 158 2 US-09-206-551-5 42 52.5 19.0 329 2 US-08-881-527A-8 44 52.5 19.0 477 1 US-08-432-016-3 | TS THEREOF | Query Match Post Some 197; DB 2; Length 855; Best Local Similarity 66.7%; Pred. No. 4.4e-19; Matches 34; Conservative 10; Mismatches 7; Indels Oy LTLGHFKEQLSKKGNYRYYFKKASDEFACGAVFEEIWDDETVLPMYEGRIL Db 798 VTLGQFKELLTKKGNYRYYFKKVSDEFDCGVVFEEVREDDTILDFFEEKII |

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Gaps
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   1 LTLGHFKEQLSKKGNYRYYFKKASDEFACGAVFEEIWDDETVLPMYEGRIL 51
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   Length 51;
   Indels
  COMPUTER: Library C. Computers: Designation of the control of the 
   APPLICANT: Constantini, Franklin
APPLICANT: Zeng, Li
TITLE OF INVENTION: AXIN GENE AND USES THEREOF
WUMBER OF SEQUENCES: 23
CORRESSPONDENCE ADDRESS:
ADDRESSEE: Cooper & Dunham LLP
   Score 190; DB 2;
Pred. No. 5.1e-18;
  1.4e-19;
  DB 2;
  9; Mismatches
  9; Mismatches
  Score 190;
Pred. No. 1
   ADDRESSEE: Cooper & Dunham LLP
STREET: 1185 Avenue of the Americas
CITY: New York
STATE: New York
COUNTRY: US
   0575/54249
  Sequence 1, Application US/08890865A
Patent No. 6307019
   NAME: White, John P
REGISTRATION NUMBER: 28,678
REFRENCE/DOCKET NUMBER: 0575
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212)278-0400
INFORMATION 10526
INFORMATION CR SEQ ID NO: 23:
SEQUENCE CHARACTERISTICS:
   ZIP: 10036
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
  68.8%;
66.7%;
   68.8%;
66.7%;
   ATTORNEY/AGENT INFORMATION:
  992 amino acids
  TELEFAX: (212)391-0526
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
   51 amino acids
  Conservative
   34; Conservative
   MOLECULE TYPE: Protein
  single
   Query Match
Best Local Similarity
Matches 34; Conserva
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  STRANDEDNESS: Sin
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  linear
  Query Match
Best Local Similarity
  GENERAL INFORMATION:
FILING DATE: 1 CLASSIFICATION:
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   US-08-890-865A-23
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  US-08-890-865A-1
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  69.2%; Score 191; DB 2; Length 900; 66.7%; Pred. No. 3.3e-18;
  CURRENT APPLICATION DATA:

REGISTRATION NUMBER: US/08/890,865A

FILING DATE: 10-JUL-1997

CLASSIFICATION: 435

ATTORNEY/AGENT INFORMATION:

NAME: White, John P

REGISTRATION NUMBER: 28,678
  SOFTWARE: Patentin Release #1.0, Version #1.30 CURRET APPLICATION DATA:
APPLICATION NUMBER: US/08/890,865A
   Sequence 23, Application US/08890865A
Patent No. 6307019
GENERAL INFORMATION:
APPLICANT: Constantini, Franklin
APPLICANT: CONSTANTION:
TITLE OF INVENTION: AXIN GENE AND USES THEREOF
NUMBER OF SEQUENCES: 23
CORRESPONDENCE ADDRESS:
  APPLICANT: Constantini, Franklin
APPLICANT: Zeng, Li
TITLE OF INVENTION: AXIN GENE AND USES THEREOF
NUMBER OF SEQUENCES: 23.
CORRESPONDENCE ADDRESS:
  Mismatches
  ADDRESSEE: Cooper & Dunham Lip
STREET: 1185 Avenue of the Americas
CITY: New York
STATE: New York
COUNTRY: US
   REFERENCE/DOCKET NUMBER: 0575/54249
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212)278-0400
TELEFAX: -(212)391-0526
   ADDRESSEE: Cooper & Dunham LLP
STREET: 1185 Avenue of the Americas
CITY: New York
STATE: New York
COUNTRY: US
  ZIP: 10036
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
CMENTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
  COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
  Sequence 4, Application US/08890865A
Patent No. 6307019
  Floppy disk
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amino acid
   TELEFAX: '(212) 391-0526
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
   Best Local Similarity 66.7
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  STRANDEDNESS: single
   MOLECULE TYPE: Protein
   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
  linear
  GENERAL INFORMATION:
  10036
  US-08-890-865A-23
  TOPOLOGY:
  US-08-890-865A-4
  COUNTRY:
   Query Match
   Matches
```

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GENERAL INFORMATION:
APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
   1 LTLGHFKEQLSKK-GNYRYYFKKASDEFACGAVFEEIWDDETVLPMYEGRIL 51
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39.7%; Score 109.5; DB 2; Length
Best Local Similarity 46.2%; Pred. No. 7.1e-07;
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   ZIP: 10036
COMPUTER READABLE FORM:
MEDLUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: 1BM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/890,865A
FILING DATE: 10-JUL-1997
CLASSIFICATION: 435
   US-08-890-865A-21
Sequence 21, Application US/08890865A
Patent No. 6307019
GENERAL INFORMATION:
APPLICANT: Constantini, Franklin
APPLICANT: Zeng, Li
TITLE OF INVENTION: AXIN GENE AND USES THEREOF;
NUMBER OF SEQUENCES: 23
CORRESPONDENCE ADDRESS:
  E: Cooper & Dunham LLP
1185 Avenue of the Americas
  US-09-949-016-10957
; Sequence 10957, Application US/09949016
; Patent No. 6812339
  ATTORNRY/AGENT INFORMATION:
NAME: White, John P
REGISTRATION NUMBER: 28,678
REFERENCE/DOCKET NUMBER: 0575
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212)278-0400
   (212)391-0526
   FORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 49 amino acids
   STRANDEDNESS: single
   MOLECULE TYPE: Protein
   TYPE: PRT
ORGANISM: Homo sapiens
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   linear
   New York
New York
   STATE: New Y COUNTRY: US
  ADDRESSEE:
  US-08-890-865A-21
   US-10-464-939-12
   TELEFAX:
SEQ ID NO 12
LENGTH: 623
  STREET:
   셤
  à
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  US-10-1949-3937-12.

US-10-1949-3937-12.

Sequence 12, Application US/10464939

Patent No. 6958238

GENERAL INFORMATION:

APPLICANT: Williams, Lewis T.

TITLE OF INVENTION: polymucleotides encoding the kinases, and methods of use

TITLE OF INVENTION: polymucleotides encoding the kinases, and methods of use

TITLE OF INVENTION: polymucleotides encoding the kinases, and methods of use

TITLE OF INVENTION: polymucleotides encoding the kinases, and methods of use

TITLE OF INVENTION: polymucleotides encoding the kinases, and methods of use

TITLE OF INVENTION: polymucleotides encoding the kinases, and methods of use

TITLE OF INVENTION: polymucleotides encoding the kinases, and methods of use

TITLE OF INVENTION: bolance 133CON

CURRENT PELLON NUMBER: 09/661,965

PRIOR FILING DATE: 2000-09-14

PRIOR FILING DATE: 1999-10-06

NUMBER OF SEQ ID NOS: 14

SOFTWARE: FastSEQ for Windows Version 4.0
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   Indels
   SOFTWARE PATENTIAN STATEM SOFTWARE BY SOFTWARE PATENTIAN BATA:

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/090,865A
FILING DATE: U-JUL-1997
CLASSIFICATION: 435
ATTONENY/AGERT INFORMATION:

NAME: White, John P
REGISTRATION NUMBER: 28,678
REGISTRATION NUMBER: 0575/54249
TELECOMMUNICATION:
TELECOMMUNICATION:
TELECOMMUNICATION:
  DB 2;
  AXIN GENE AND USES THEREOF
   Query Match 39.7%; Score 109.5; DB 2
Best Local Similarity 46.2%; Pred. No. 3.3e-08;
Matches 24; Conservative 10; Mismatches 15
   E: Cooper & Dunham LLP
1185 Avenue of the Americas
   MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
   Sequence 22, Application US/08890865A Patent No. 6307019 GENERAL INFORMATION:
  APPLICANT: Constantini, Franklin APPLICANT: Zeng, Li TITLE OF INVENTION: AXIN GENE ANI UMBER OF SEQUENCES: 23 CORRESPONDENCE ADDRESS: ADDRESSE COOPER & Dunham LLP
   INFORMATION FOR SEQ ID NO: 22:
   (212)391-0526
  LENGTH: 50 amino acids
  SEQUENCE CHARACTERISTICS:
  TOPOLOGY: linear
MOLECULE TYPE: Protein
   TYPE: amino acid
STRANDEDNESS: single
  ZIP: 10036
COMPUTER READABLE FORM:
  CITY: New York
STATE: New York
COUNTRY: US
   US-08-890-865A-22
   US-08-890-865A-22
   US-10-464-939-12
  STREET:
                            유
   ò
  셤
```

Gaps

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GENERAL INFORMATION:

GENERAL INFORMATION:

APPLICANT: Homburger et al.

TITLE OF INVENTION: Nucleic acids and proteins of Drosophila melanogaster FILE REFERENCE: File Reference: 7326-094

CURRENT APPLICATION NUMBER: US/09/270,767

CURRENT FILING DATE: 1999-03-17

NUMBER OF SEQ ID NOS: 62517

SOFTWARE: Patentin Ver. 2.0

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; Sequence 43189, Application US/09270767
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OPERATING SYSTEM: PC-DOS/MS-DOS
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; Sequence 42, Application US/09398395A
; Patent No. 6468772
; GENERAL INFORMATION:
   No. 64687721, Joseph P. Starks, Courtney M. Manna, Kathleen R.
  , ORGANISM: Drosophila melanogaster
US-09-270-767-43189
  ATTORNEY/AGENT INFORMATION:
NAME: White, John P
REGISTRATION NUMBER: 0678
REFERENCE/DOCKET NUMBER: 0579
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 278-0400
TELEPHONE: (212) 391-0526
  28,678
  INFORMATION FOR SEQ ID NO: 20: SEQUENCE CHARACTERISTICS:
  Match 34.4%;
Local Similarity 39.6%;
les 21; Conservative 1
  51 amino acids
   TYPE: amino acid
STRANDEDNESS: single
   TOPOLOGY: linear
MOLECULE TYPE: Protein
                                  ZIP: 10036
COMPUTER READABLE FORM:
New York
   APPLICANT: Chappell, APPLICANT: No. 6468
   US-08-890-865A-20
                  COUNTRY:
  APPLICANT:
  TYPE: PRT
   APPLICANT:
   Query Match
   Matches
   a
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  GENERAL INFORMATION:
APPLICANT: VENTER, J. Craig et al.
APPLICANT: VENTER, J. Craig et al.
APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
FILE REFERENCE: CLOO1307
CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-10-0-0
PRIOR APPLICATION NUMBER: 60/241,755
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PRIOR FILING DATE: 2000-10-03
PRIOR FILING DATE: 2000-09-08
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   37.0%; Score 102; DB 2; Length 738; 41.2%; Pred. No. 1e-05; tive 8; Mismatches 20; Indels
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41.2%; Pred. No. 8.1e-05;
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             CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
PRIOR PELICATION NUMBER: 60/241,755
PRIOR PELING DATE: 2000-10-20
PRIOR PELING DATE: 2000-10-30
PRIOR PELING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR PELING DATE: 2000-09-08
PRIOR FILING DATE: 2000-09-08
PRIOR FILING DATE: 2000-09-08
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SPIOR FILING DATE: 2000-09-08
   3: Cooper & Dunham LLP
1185 Avenue of the Americas
   Sequence 20. Application US/08890865A
Patent No. 6307019
GENERAL INFORMATION:
APPLICANT: Constantini, Franklin
APPLICANT: Zeng, Li
TITLE OF INVENTION:
NUMBER OF SEQUENCES: 23
CORRESPONDENCE ADDRESS:
ADDRESSEE: COOPER & Dunham LLP
STREET: 1185 Avenue of the America
  Sequence 6495, Application US/09949016
Patent No. 6812339
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US-08-890-865A-20
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Gaps

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Length 601;

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  22.1%; Score 61; DB 29.2%; Pred. No. 5; Live 12; Mismatches
                                      CURRENT FILING DATE: 2001-06-29
PRIOR APPLICATION NUMBER: 09/398,395
PRIOR FILING DATE: 1999-09-17
PRIOR APLICATION NUMBER: 60/100,993
PRIOR FILING DATE: 1998-09-18
PRIOR FILING DATE: 1998-09-18
PRIOR FILING DATE: 1999-09-18
PRIOR FILING DATE: 1999-09-22
PRIOR FILING DATE: 1999-08-23
NUMBER OF EXQ ID NOS: 58
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CURRENT APPLICATION NUMBER: US/09/895,752
CURRENT FILING DATE: 2001-06.29
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Matches 14; Conservative
  ORGANISM: Ricinus communis
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Best Local Similarity
Matches 14; Conserv
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US-09-903-012B-42
  US-09-895-752-42
  LENGTH: 601
   SEQ ID NO 42
  SOFTWARE: F
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  TYPE: PRT
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  ; Pred. No. 5;
12; Mismatches
  22.1%; Score 61; DB 2
29.2%; Pred. No. 5;
tive 12; Mismatches
   APPLICANT: Chappell, Joseph
APPLICANT: No. 64953541, Joseph
APPLICANT: No. 64953541, Joseph
APPLICANT: STARKS, COURTERY M.
APPLICANT: STARKS, COURTERY M.
APPLICANT: Manna, Kathleen R.
TITLE OF INVENTION: SYNTHASES
FILE REFERENCE: 07678-025011
CURRENT FILING DATE: 2001-06-22
PRIOR APPLICATION NUMBER: 09/398,395
PRIOR APPLICATION NUMBER: 60/130,628
PRIOR APPLICATION NUMBER: 60/130,628
PRIOR APPLICATION NUMBER: 60/130,628
PRIOR APPLICATION NUMBER: 60/130,628
PRIOR APPLICATION NUMBER: 60/150,262
PRIOR APPLICATION NUMBER: 60/150,262
PRIOR FILING DATE: 1999-08-23
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SOFTWARE: FASTSEQ for Windows Version 3.0
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TITLE OF INVENTION: SYNTHASES
FILE REFERENCE: 07678-025001
CURRENT PELING DATE: 1999-09-17
PRIOR APPLICATION NUMBER: 60/100,993
PRIOR FILING DATE: 1999-09-18
PRIOR FILING DATE: 1999-04-22
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PRIOR FILING DATE: 1999-08-23
NUMBER OF SEQ ID NOS: 58
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Pred. No.
  US-09-887-586A-42; Sequence 42, Application US/09887586A; Patent No. 6495354
  Sequence 42, Application US/09895752
Patent No. 6559297
  GENERAL INFORMATION:
APPLICANT: Chappell, Joseph
APPLICANT: No. 65592971, Joseph P.
APPLICANT: No. 65592971, Money M.
APPLICANT: Manna, Kathleen R.
TITLE OF INVENTION: SYNTHASES
  Query Match
22.1%;
Best Local Similarity 29.2%;
Matches 14; Conservative 1:
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ORGANISM: Ricinus communis
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  US-09-398-395A-42
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US-09-895-752-42
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   DB 2;
   Score 61; DB 2
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  APPLICANT: Chappell, Joseph
APPLICANT: Chappell, Joseph
APPLICANT: Starks, Courtney M.
APPLICANT: Starks, Courtney M.
APPLICANT: Manna, Kathleen R.
TITLE OF INVENTION: SYNTHASES
FILE REFERENCE: 07678-02501
CURRENT APPLICATION NUMBER: US/09/903,012B
CURRENT FILING DATE: 12091-07-11
PRIOR APPLICATION NUMBER: 60/100,993
PRIOR FILING DATE: 1999-04-22
PRIOR FILING DATE: 1999-04-22
PRIOR APPLICATION NUMBER: 60/130,628
PRIOR FILING DATE: 1999-04-22
PRIOR APPLICATION NUMBER: 60/150,262
PRIOR FILING DATE: 1999-08-23
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Sequence 42, Application US/09903012B
Patent No. 6569656
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US-09-903-012B-42
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300, App
12, Appl
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Sequence 8, Appli
  April 20, 2006, 15:57:53 ; Search time 17.7177 Seconds (without alignments) 1202.714 Million cell updates/sec
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GenCore version 5.1.7.
Copyright (c) 1993 - 2006 Biocceleration Ltd.
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US-10-73-860-1797

US-10-78-813-81

US-10-786-720-36

US-10-786-720-35

US-10-786-720-35

US-10-182-936A-91

US-10-477-238A-670

US-10-680-287A-670

US-10-680-287A-670

US-10-092-900A-270

US-10-094-749-2273

US-10-094-749-2273

US-10-094-749-2273

US-10-094-749-2273

US-10-097-143-2028

US-10-678-639-41

US-10-678-639-41
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Maximum Match 100%
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Maximum DB seq length: 200000000
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## ALIGNMENTS

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TITLE OF INVENTION: Colon and Colon Cancer Associated Polynucleotides and Polypepti TITLE OF INVENTION: Colon and Colon Cancer Associated Polynucleotides and Polypepti CURRENT APPLICATION NUMBER: US/10/106,698
CURRENT FILING DATE: 2002-03-27
PRIOR APPLICATION NUMBER: US 60/157,137
PRIOR PILING DATE: 1999-09-29
PRIOR PILING DATE: 1999-09-29
PRIOR PILING DATE: 1999-11-03
NUMBER OF SEC ID NOS: 8564
SOFTWARE: Patentin Ver: 3.0
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NAME/KEY: MISC_FEATURE
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  any of the naturally occurring L-amino acids
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Publication No. US20030109690A1
GENERAL INFORMATION:
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OTHER INFORMATION: Xaa equals
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   ORGANISM: Homo sapiens
  NAME/KEY: MISC_FEATURE
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Matches 50; Conserv
  LOCATION:
  Query Match
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APPLICANT: O'Toole, Margot
APPLICANT: Liu, Wei
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR DIAGNOSING AND TREATING AUTOIMMUNE
TITLE OF INVENTION: DISEASES
                                 TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR DIAGNOSING AND TREATING COLON TITLE OF INVENTION: CANCERS FILE REFERENCE: AM100927 (031996-002000)
CURRENT APPLICATION NUMBER: US/10/751,736
CURRENT FILING DATE: 2003-01-06
PRIOR PILING DATE: 2003-01-06
NUMBER OF SEQ ID NOS: 54873
SOFTWARE: PATENTIN VESTOR 3.2
  APPLICANT: KLEIN, Peter S.

TITLE OF INVENTION: AXIN DOMAIN-LIKE POLYPEPTIDE INHIBITORS OF GLYCOGEN
TITLE OF INVENTION: SYNTHASE KINASE 3 (BETA) ACTIVITY AND ACTIVATORS OF WNT
TITLE OF INVENTION: SIGNALING
FILE REFERENCE: 209596.0391/30601
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CURRENT FILING DATE: 2004-02-26
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Publication No. US20040191818A1
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APPLICANT: Wyeth
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Patent No. US20010052137A1
GENERAL INFORMATION:
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US-10-264-049-2846
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  APPLICANT: Aziz, Natasha
APPLICANT: Ginsburg, Wendy M.
APPLICANT: Ginsburg, Wendy M.
APPLICANT: Zlotnik, Albert
TITLE OF INVENTION: Methods of Diagnosis of Soft Tissue Sarcoma, Compositions
TITLE OF INVENTION: Methods for Screening for Soft Tissue Sarcoma Modulators
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  Score 272; DB 4; Length 347;
Pred. No. 1.9e-28;
APPLICANT: Birse et al.
TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies FILE REFERENCE: PA137191
CURRENT APPLICATION NUMBER: US/10/264,049
CURRENT FILING DATE: 2002-10-04
PRIOR APPLICATION NUMBER: PCT/US01/18569
PRIOR FILING DATE: US 60/209,467
PRIOR FILING DATE: 2000-06-07
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US-10-723-860-1797
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LENGTH: 843
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TITLE OF INVENTION: Reagents and Method for Modulating DKK-Mediated Interactions FILE REFERENCE: 032796-143
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  ö
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  Gaps
   APPLICANT: Babij, Philip
APPLICANT: Yaworsky, Paul
APPLICANT: Bex, Frederick J. III
APPLICANT: Bodine, Peter Van Neet
TITLE OF INVENTION: Transgenic Animal Model of Bone Mass Modulation
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PRIOR PILING DATE: 2002-02-01
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PRIOR PILING DATE: 2002-03-04
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CURRENT FILING DATE: 2003-11-10
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PRIOR FILING DATE: 2001-05-11
PRIOR PELICATION NUMBER: US 60/291,311
PRIOR FILING DATE: 2001-05-17
PRIOR FILING DATE: 2001-05-17
PRIOR APPLICATION NUMBER: US 60/353,058
PRIOR FILING DATE: 2002-02-01
PRIOR PELING DATE: 2002-02-01
PRIOR FILING DATE: 2002-03-04
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SOFTWARE: FastSEQ for Windows Version 4.0
   Sequence 670, Application US/10477238A Publication No. US20040221326A1 GENERAL INFORMATION:
   US-10-182-936A-91

Sequence 91, Application US/10182936A

Publication No. US20040038860A1

GENERAL INFORMATION:
APPLICANT: Anlen, Kristina M.
APPLICANT: Anlsowicz, Anthony
APPLICANT: Bhat, Bheem
   Damagnez, Veronique
Robinson, John
Yaworsky, Paul
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   APPLICANT: O'Toole, Margot
APPLICANT: Liu, Wei
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR DIAGNOSING AND TREATING AUTOIMMUNE
TITLE OF INVENTION: DISEASES.
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   APPLICANT: John P. Carulli et al.
TITLE OF INVENTION: THE HIGH BONE MASS GENE OF 11q13.3
FILE REFERENCE: 022796-021
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Score 191;

69.2%;

Query Match

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   APPLICANT: Liu, Wei
TITLE OF INVENTION: HBM Variants that Modulate Bone Mass and Lipid Levels
FILE REFERENCE: 032796-135
CURRENT APPLICATION NUMBER: US/10/477,173
CURRENT FILING DATE: 2003-11-10
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   APPLICANT: Yaworsky, Paul
APPLICANT: Bex, Frederick J: III
APPLICANT: Bodine, Peter Van Nest
TITLE OF INVENTION: Transpenic Animal Model of Bone Mass Modulation
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66.7%; Pred. No. 6.7e-17;
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PRIOR APPLICATION NUMBER: US 60/291,311
PRIOR FILING DATE: 2001-05-17
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  Yaworsky, Paul
Morales, Arturo J.
Graham, James R.
Anisowicz, Anthony
   Allen, Kristina M.
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APPLICANT: Alsobrook, John P.
PPPLICANT: Lepley, Denise M.
APPLICANT: Rieger, Daniel K.
APPLICANT: Rieger, Daniel K.
APPLICANT: Burgess, Catherine E.
ITLE OF INVENTION: No. US20040043382Alel Proteins and Nucleic Acids Encoding Same
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   PRIOR APPLICATION NUMBER: USSN 60/274,322
   FILING DATE: 2001-03-08
APPLICATION NUMBER: USSN 60/325,681
  Sequence 270, Application US/10092900A Publication No. US20040043382A1 GENERAL INFORMATION:
   APPLICANT: Padigaru, Muralidhara
APPLICANT: Spytek, Kimberly A.
APPLICANT: Shenoy, Suresh G.
APPLICANT: Taupier Jr., Raymond J.
APPLICANT: Pena, Carol B.A.
APPLICANT: Li, Li
   Tchernev,, Velizar T
Fernandes, Elma R.
   Corine A.M
  Zerhusen, Bryan D.
Gusev, Vladimir Y.
Ji, Weizhen
   Gorman, Linda
Miller, Charles E.
  Guo, Xiaojia Sasha
  Kekuda, Ramesh
Patturajan, Meera
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  Anderson, David W
  FILING DATE: 2001-03-08
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Gerlach, Valerie
   Casman, Stacie J
  Catterton, Elina
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  Leite, Mario W.
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PRIOR PILING DATE: 2001-05-31
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   APPLICANT: MASUHO, YASUHIKO
TITLE OF INVENTION: NOVEL FULL-LENGTH CDNA
FILE REFERENCE: 084335/0160
CURRENT APPLICATION NUMBER: US/10/094,749
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  PRIOR APPLICATION NUMBER: JP 2001-328381
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  SUGIYANA, TOMOYASU
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ISHII, SHIZUKO
YAMAMOTO, JUN-ICHI
ISONO, YUUKO
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  YOSHIKAWA, TSUTOMU
OTSUKA, MOTOYUKI
  AMECHIKA, ICHIRO
  NAGAHARI, KENJI
   OTSUKA, KAORU
NAGAI, KEIICHI
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SOFTWARE: Patentin Ver. 2.
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   SEKI, NAOHIKO
   ORGANISM: Homo sapiens
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  APPLICANT: YOSHIKAWA, TSUTOMU
APPLICANT: OTSUKA, MOTOYUKI
APPLICANT: NAGAHARI, KENJI
APPLICANT: MASUHO, YASUHIKO
TITLE OF INVENTION: NOVEL FULL-LENGTH CDNA
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ISONO, YUUKO
   IRIE, RYOTARO
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Copyright (c) 1993 - 2006 Biocceleration Ltd
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US-11-095-561-851
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## ALIGNMENTS

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APPLICANT: Alexandrov, Nickolai et al.
APPLICANT: Alexandrov, Nickolai et al.
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; APPLICANT: Abad, Mark S. et al.
; TITLE OF INVENTION: Genes and Uses for Plant Improvement;
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  APPLICANT: Rabinowitz, Joshua D.
APPLICANT: Rabinowitz, Joshua D.
APPLICANT: Schwaizer, Johannas
APPLICANT: Schwaizer, Johannas
APPLICANT: Carrick, Deanna Maxie
APPLICANT: Arbor Vita Corporation
TITLE OF INVENTION: MODECULAR INTERACTION IN MODECULAR INTERACTION IN MODECULAR INTERACTION IN WARENT US/10/485, 788A
CURRENT FILING DATE: 2004-02-03
PRIOR FILING DATE: 2001-08-03
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PRIOR APPLICATION NUMBER: US 60/360,061
PRIOR APPLICATION NUMBER: WO PCT/US02/24655
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  APPLICANT: Lu, Peter S.
APPLICANT: Lu, Peter S.
APPLICANT: Rabinowitz, Joshua D.
APPLICANT: Schwicze, Johannes
APPLICANT: Carrick, Deanna Marie
APPLICANT: Arbor Vita Corporation
TITLE OF INVENTION: Molecular Interactions in Cells
FILE REFERENCE: 20054-003320US
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PRIOR FILING DATE: 2001-08-03
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PRIOR FILING DATE: 2002-02-25
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TYPE: PRT
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Fublication No. US20060073161A1
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  APPLICANT: Hirsch, Joel
TITLE OF INVENTION: VOLTAGE-DEPENDENT CALCIUM CHANNEL BETA SUBUNIT FUNCTIONAL CORE
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TITLE OF INVENTION: Genes and Uses for Plant Improvement
FILE REFERENCE: 38-21(53450)B EP
CURRENT APPLICATION NUMBER: US/11/087,099
CURRENT FILING DATE: 2005-03-22
NUMBER OF SEQ ID NOS: 12464
SEQ ID NO 589
  146 KQDTKKGNLRYYPYNINWNY---GLFPQTWEDPTL 177
   S1 KQDTKKGNLRYYPYNINWNY---GLFPQTWEDPTL 82
  8 EQLSKKGNYRYYFKKASDEFACGAVFEEIWDDETV 42
  ; LOCATION: (1)..(315)
; OTHER INFORMATION: unsure at all Xaa locations US-11-087-099-589
   8 EQLSKKGNYRYYFKKASDEFACGAVFEEIWDDETV
    Best Local Similarity 34.3%; Pred. No. 5.9; Matches 12; Conservative 8; Mismatches
   Score 55; DB '
Pred, No. 16;
5; Mismatches
  232 RIYFESAFNRFDCIVIAGSIFEVVW 256
  17 RYYFKKASDEFAC----GAVFEEIW 37
   US-11-087-099-589
Sequence 589, Application US/11087099
Publication No. US20060041961A1
GENERAL INFORMATION:
   Sequence 35, Application US/11126313
Publication No. US20050288489A1
GENERAL INFORMATION:
  US-11-126-313-36
; Sequence 36, Application US/11126313
; Publication No. US20050288489A1
; GENERAL INFORMATION:
   Query Match
Best Local Similarity 40.0%;
Matches 10; Conservative
  ORGANISM: Anopheles gambiae US-11-126-313-35
   ORGANISM: Oryza sativa
  NAME/KEY: unsure
  RESULT 8
US-11-126-313-35
   SEQ ID NO 35
  LENGTH:
```

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Sequence 27905, Application US/11096568A
Publication No. US20060048240A1
GENERAL INPORMATION:
APPLICANT: Alexandrov, Nickolai et al.
TITLE OF INVENTION: Sequence-Determined DNA Fragments and Corresponding Polypeptide
TITLE OF INVENTION: Therby
   Length 615;
   DB 7; Length 699;
   ----FEGKKL 428
   14; Indels
  6 FKEQLSKKGNYRYYFKKASDEFACGAVFEEIWDDETVLPMYEGRIL
  6 FKEQLSKKGNYRYYFKKASDEFACGAVFEEIWDDETVLPMYEGRIL
  DB 7;
   Score 53.5; DE Pred. No. 39; 5; Mismatches
   NAME/KEY: misc_feature
| LOCATION: (1)..(615)
| OTHER INFORMATION: Ceres Seq. ID no. 2133277
US-11-096-5688-27906
  NAME/KEY: misc_feature
| LOCATION: (1)...(699)
| HTER INDORMATION: Ceres Seq. ID no. 2133276
US-11-096-568A-27905
   Score 53.5; D
Pred. No. 45;
5; Mismatches
   FILE REFERENCE: 2750-1592PUS2
CURRENT APPLICATION NUMBER: US/11/096,568A
CURRENT FILING DATE: 2005-04-01
NUMBER OF SEQ ID NOS: 34471
SEQ ID NO 27905
   478 FLEKLKKKGIEVLYMVDAIDEYAIGQLKE---
   394 FLEKLKKKGIEVLYMVDAIDEYAIGQLKE
   Search completed: April 20, 2006, 16:07:54 Job time : 3.82586 secs
   TYPE: PRT
ORGANISM: Arabidopsis thaliana
   19.4%;
34.8%;
   Query Match
Best Local Similarity 34.8%;
Matches 16; Conservative
   Best Local Similarity 34.8
Matches 16; Conservative
   TYPE: PRT
ORGANISM: Arabidopsis
CURRENT FILING DATE:
NUMBER OF SEQ ID NOS:
SEQ ID NO 27906
   JS-11-096-568A-27905
  Query Match
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   TITLE OF INVENTION: Sequence-Determined DNA Fragments and Corresponding Polypeptides TITLE OF INVENTION: Therby
TITLE OF INVENTION: Therby
CURRENT APPLICATION NUMBER: US/11/096,568A
CURRENT FILING DATE: 2005-04-01
NUMBER OF SEQ ID NOS: 34471
LENGTH: 594
  TITLE OF INVENTION: Sequence-Determined DNA Fragments and Corresponding Polypeptides IITLE OF INVENTION: Therby
   APPLICANT: CARGILL, Michele et al.
TITLE OF INVENTION: GENETIC POLYMORPHISMS ASSOCIATED WITH
TITLE OF INVENTION: CARDIOVASCULAR DISORDERS AND DRUG RESPONSE, METHODS OF
TITLE OF INVENTION: DETECTION AND USES THEREOF
FILE REFERENCE: CLOOISS9
CURRENT APPLICATION NUMBER: US/10/995,561
CURRENT PILING DATE: 2004-11-24
NUMBER OF SEQ ID NOS: 85702
SOFTWARE: FABELSEQ for Windows Version 4.0
  Gaps
   2;
   DB 7; Length 594;
  6 FKEQLSKKGNYRYYFKKASDEFACGAVFBEIWDDETVLPMYEGRIL 51
    374 RDSLDKHRTDKREFLYTRDQLECGATHDELWNAAQLEMVHVGKM 417
   17; Indels
   1 LTLGHFKEQLSKKGNYRYYFKKASDEFACGAVFEEIWDD 39
  852 VSFSQIKSKLIRVENFEAYFKKQQADSNCG--FAEEYED
  DB 6;
  8; Mismatches
   NAME/KEY: misc_feature
LOCATION: (1)..(594)
CTHER INVERMATION: Ceres Seq. ID no. 2133278
US-11-096-568A-27907
   5; Mismatches
  19.4%; Score 53.5; 34.8%; Pred. No. 37;
  19.6%; Score 54; DB 30.8%; Pred. No. 70;
  FILE REFERENCE: 2750-1592PUS2
CURRENT APPLICATION NUMBER: US/11/096,568A
  Sequence 27907, Application US/11096568A Publication No. US20060048240A1 GENERAL INFORMATION:
  Sequence 27906, Application US/11096568A Publication No. US20060048240A1 GENERAL INFORMATION:
  APPLICANT: Alexandrov, Nickolai et al
  Sequence 851, Application US/10995561 Publication No. US20050272054A1 GENERAL INFORMATION:
   APPLICANT: Alexandrov, Nickolai
  ORGANISM: Arabidopsis thaliana
   Query Match
Best Local Similarity 34.8<sup>†</sup>
Matches 16; Conservative
  Conservative
  ORGANISM: Homo sapiens
US-10-995-561-851
  Query Match
Best Local Similarity
Matches 12; Conserv
   RESULT 13
US-11-096-568A-27907
   SEQ ID NO 851
LENGTH: 1178
   TYPE: PRT
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